Machine learning Part 2 Model selection & validation

Jérôme Dockès & Nikhil Bhagwat

MAIN tutorials course 2021-11-26

Outline

Introduction: cross-validation

Model and hyperparameter selection

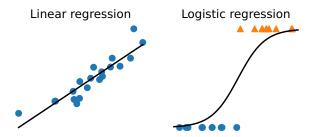
Dimensionality reduction

Conclusion

Recap of part 1

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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Supervised learning

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- · Classification: logistic regression

Regularization

• ℓ_2 a.k.a. ridge regularization

Model evaluation and selection

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

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

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- $Y \in \mathbb{R}$: output (a.k.a. target, dependent variable) to predict
- $X \in \mathbb{R}^p$: features (a.k.a. inputs, regressors, descriptors, independent variables)
- $E \in \mathbb{R}$: unmodelled noise
- f: the function we try to approximate

Example: linear regression

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

$$= \beta_0 + \sum_{j=1}^{p} X_j \beta_j + E$$
(2)
(3)

$$j=1$$
"learning" = estimating $\beta_{\alpha} \in \mathbb{R}$ and $\beta \in \mathbb{R}^p$

(1)

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:

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

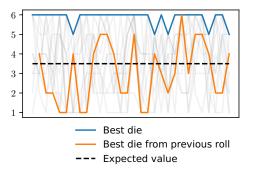
$$= \sum_{i=1}^{n} (\mathbf{y}_{i} - \hat{\boldsymbol{\beta}}_{0} - \sum_{j=1}^{p} \mathbf{X}_{ij}\,\hat{\boldsymbol{\beta}}_{j})^{2}$$
(5)

"Fitting" the parameters to X, y.

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



Estimating prediction performance

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.

scikit-learn "estimator API": fit; predict

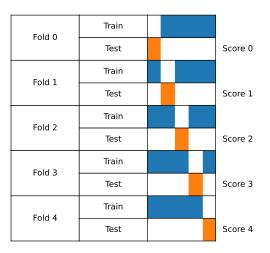
```
estimator = Ridge()
estimator.fit(X_train, y_train)
predictions = estimator.predict(X_test)
```

```
https://scikit-learn.org/stable/getting_started.html
sklearn.linear_model.Ridge
```

Evaluating performance with sklearn.metrics

```
estimator = Ridge()
estimator.fit(X_train, y_train)
predictions = estimator.predict(X_test)
mse = metrics.mean_squared_error(y_test, predictions)
https://scikit-learn.org/stable/getting_started.html
sklearn.linear_model.Ridge
sklearn.metrics
more info on model evaluation
ex_01_fit_predict.py
```

Cross-validation



scikit-learn.org/stable/modules/cross_validation.html
sklearn.model_selection.cross_validate
ex_02_cross_validate.py

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Introduction: cross-validation

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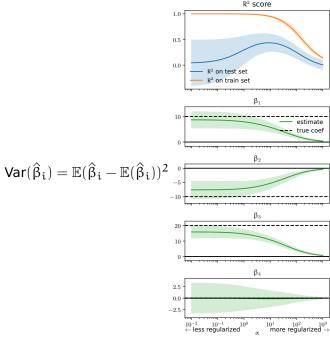
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Regularization

Example: ridge regression

$$\underset{\beta,\beta_0}{\operatorname{argmin}} \|\mathbf{y} - \beta_0 - \mathbf{X} \, \boldsymbol{\beta}\| + \alpha \, \|\boldsymbol{\beta}\|_2^2 \tag{6}$$



 $\mathsf{Bias}(\hat{\beta}_{i}) = \mathbb{E}(\hat{\beta}_{i}) - \beta_{i}$

Setting hyperparameters

How can we choose the ridge hyperparameter α ?

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

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

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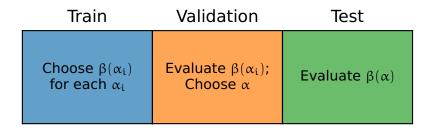
When you hear "best", "maximum", "select", ... think "bias" Setting the parameters

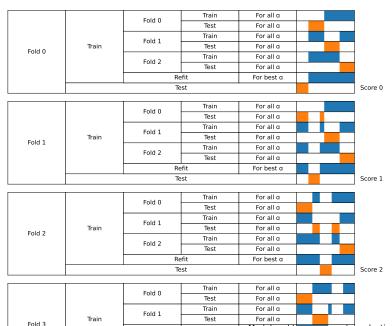
- Select β that gives the **best** prediction on training data
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Setting the hyperparameters

- Repeat step 1 for a few values of α , 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





Train

FALL 2

Medeliand hyperparameter selection

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Implementing nested CV

 ${\tt ex_05_nested_cross_validation.py}$

Outline

Introduction: cross-validation

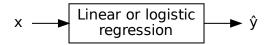
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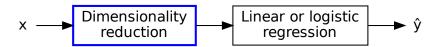
Conclusion

Dimensionality reduction

Until now



Add a step in the pipeline: simplifying the inputs



Dimensionality reduction

$$\hat{y} = \hat{\beta}_0 + \hat{\beta}_1 X_{:,1} + \hat{\beta}_2 X_{:,1} + \dots + \hat{\beta}_p X_{:,p}$$
 (7)

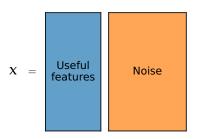
Problems when the number of features p becomes large

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

Simulated data for linear regression

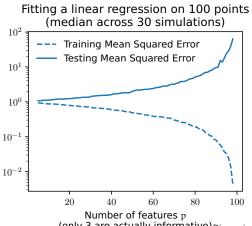
- Generate $X \in \mathbb{R}^{n \times 3}$, $\beta \in \mathbb{R}^3$, $e \in \mathbb{R}^n$ and $y = X\beta + e \in \mathbb{R}^n$
- Append columns containing random noise to X
- Now $X \in \mathbb{R}^{n \times p}$, with $p \geqslant 3$, but only the first 3 columns are linked with y
- Split into training 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 p can fit exactly (0 training error) any set of p+1 points.
- Risk of overfitting: fitting exactly training data but failing on test data



Univariate feature selection

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

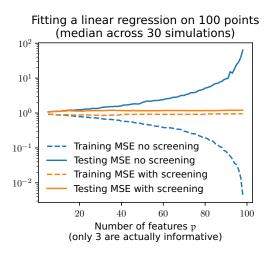
Simple (linear) association criteria

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

Read more in the scikit-learn user guide scikit-learn feature selection

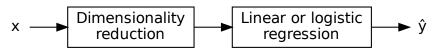
Univariate feature selection

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

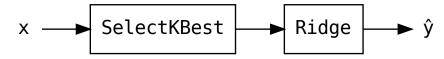


Dataset transformations

Typical pipeline



Example



scikit-learn "transformer API": fit; transform

```
transformer = SelectKBest()
transformer.fit(X_train)
transformed_X = transformer.transform(X_train)
```

can also be written:

```
transformer = SelectKBest()
transformed_X = transformer.fit_transform(X_train)
```

scikit-learn feature selection

feature_selection.SelectKBest

fit:

- compute ANOVA or correlation for each column of X
- Remember the indices of the k columns with highest scores

transform:

• Index input to keep only the k selected columns

sklearn.feature_selection.SelectKBest

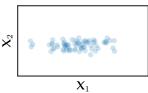
scikit-learn "transformer API": fit; transform

```
transformer = SelectKBest()
transformed_X_train = transformer.fit_transform(X_train)
transformed_X_test = transformer.transform(X_test)
```

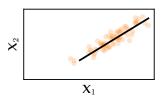
ex_04_feature_selection.py

Solution 2: linear decomposition methods

Maybe OK to drop X_2 :



Data low-dimensional but no feature can be dropped:

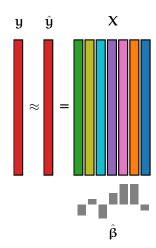


Find a better referential in which to represent the data

Linear regression: projection on the column

space of X

$$\hat{\mathbf{y}} = \mathbf{X} \, \hat{\mathbf{\beta}}$$
 (8)

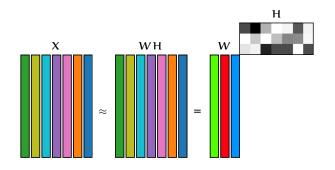


- Too many features: high variance & unstable solution
- Feature selection: drop some columns of X
- Other ways to build a family of k vectors on which to regress y?

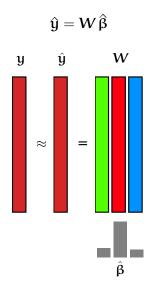
Linear decomposition: low-rank approximation of \mathbf{X}

Minimize

$$\|X - WH\|_{F}^{2} = \sum_{i,j} (X_{i,j} - (WH)_{i,j})^{2}$$
 (9)



Linear regression after dimensionality reduction



(10)

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 $\hat{\beta}$

$$x \in \mathbb{R}^p o \mathsf{projection} o w \in \mathbb{R}^k o \langle \cdot \,,\, \hat{eta}
angle o \hat{y} \in \mathbb{R}$$
 (11)

Principal Component Analysis

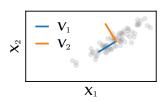
Singular Value Decomposition of X:

$$X = \mathbf{U} \, \mathbf{S} \, \mathbf{V}^{\mathsf{T}} \tag{12}$$

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)
- $S \succeq 0$ 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

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

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

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

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

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

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

$$\mathbf{v}_{5}$$

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

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

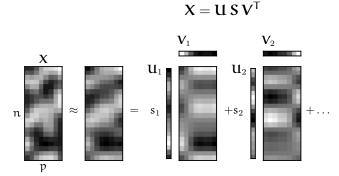
$$\mathbf{V}^\mathsf{T}\,\mathbf{V} = \mathrm{I}_{\mathrm{p}}$$

$$\mathbf{V}^\mathsf{T} \mathbf{V} = \mathbf{I}_{\mathbf{p}}$$

(14)

(15)

Singular Value Decomposition



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

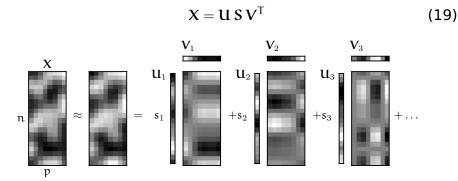
$$\mathbf{V}^{\mathsf{T}} \, \mathbf{V} = \mathrm{I}_{\mathrm{p}} \tag{18}$$

$$\mathbf{V}^\mathsf{T} \mathbf{V} = \mathbf{I}_{\mathfrak{p}}$$

(18)

(16)

Singular Value Decomposition



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

$$\mathbf{V}^{\mathsf{T}} \, \mathbf{V} = \mathrm{I}_{\mathrm{p}} \tag{21}$$

$$\mathbf{V}^{\mathsf{T}}\,\mathbf{V} = \mathbf{I}_{\mathsf{p}} \tag{21}$$

Other decomposition methods

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

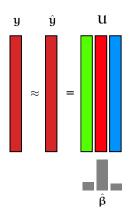
- · Dictionary Learning
- Non-negative Matrix Factorization
- · K-means clustering
- ...

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 $s_j^2/(s_j^2+\alpha)$
- PC regression sets the coordinates to 0 except for those corresponding to the k largest s_j: shrinks by a factor 1_{j≤k}



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Some pitfalls with cross-validation

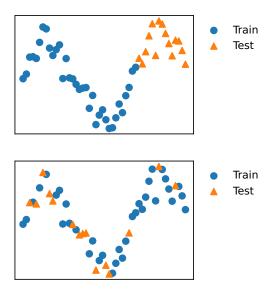
- Overfitting the hyperparameters
 - select hyperparameters with nested CV sklearn.model_selection.GridSearchCV

Fitting part of the pipeline on the whole dataset

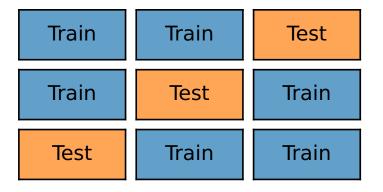
- use sklearn.pipeline.Pipeline
- Ignoring dependencies between samples
 - e.g. time series: use appropriate cross-validation iterator
- Ignoring dependencies between CV scores
 - Training sets overlap: cross-validation scores of different splits are not independent
- Over-interpreting good CV scores
 - Good CV scores do not mean the model will always perform well on a new dataset

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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.