# **Conformal Prediction**

A Tutorial on Predicting with Confidence

Henrik Linusson<sup>1</sup>, Ulf Johansson<sup>2</sup>, Tuve Löfström<sup>1</sup>, Henrik Boström<sup>3</sup>, Alex Gammerman<sup>4</sup> August 12, 2018

<sup>1</sup>University of Borås, Sweden. Email: {henrik.linusson, tuve.lofstrom}@hb.se <sup>2</sup>Jönköping University, Sweden. Email: ulf.johansson@ju.se <sup>3</sup>KTH, Royal Institute of Technology, Sweden. Email: henrik.bostrom@dsv.su.se <sup>4</sup>Royal Holloway, University of London, United Kingdom. Email: a.gammerman@cs.rhul.ac.uk

## Agenda

- Purpose and goal
- A motivating example
- Conformal prediction at a glance
- Conformal classification
- Conformal regression
- Validity and efficiency
- Considerations and modifications
- Conformal classification a critical look
- Venn predictors
- Nonconformist conformal prediction in Python
- Other scenarios and suggested reading

# Purpose and goal

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- There is absolutely no magic involved only mathematics!
- Hot topic recently picked up by both academia and industry
- Plenty of open questions, i.e., research opportunities

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- ...or even be part of the small but growing conformal society
- Disclaimer: I come from machine learning not algorithmic theory...

A motivating example

## How good is your prediction?

You want to estimate the risk of cancer recurrence in patient  $x_{k+1}$ 

To your disposal, you have:

- 1. A set of historical observations  $(x_1, y_1), \ldots, (x_k, y_k)$ 
  - x<sub>i</sub> describes a patient by age, tumor size, etc
  - $y_i$  is a measurement of cancer recurrence in patient  $x_i$
- 2. Some machine learning (classification or regression) algorithm

```
import pandas as pd
```

breast\_cancer = pd.read\_csv('./data/breast-cancer.csv')

```
# (x_1, y_1), ..., (x_k, y_k)
x_train = breast_cancer.values[:-1, :-1]
y_train = breast_cancer.values[:-1, -1]
```

```
# (x_k+1, y_k+1)
x_test = breast_cancer.values[-1, :-1]
y_test = breast_cancer.values[-1, -1]
```

from sklearn.neighbors import KNeighborsClassifier

```
knn = KNeighborsClassifier(n_neighbors=5)
knn.fit(x_train, y_train)
```

```
print(knn.predict(x_test))
print(knn.predict_proba(x_test))
```

```
['no-recurrence-events']
[[ 0.8 0.2 ]]
```

## How good is your prediction, really?

- Your classifier says that the patient will have no recurrence events. Is it right?
- Your probability estimator says it's 80% likely that the patient won't have a recurrence event.
  - How good is the estimate?
- Your regression model says the patient should have 0.4 recurrence events in the future.
  - How close is that to the true value?

Will you trust your model?

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We expect past performance to indicate future performance.

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- The model is 71% accurate on the test data, so we assume it's accurate for 71% of production data.
- The model has an AUC of 0.65 on the test data, so we assume it has an AUC of 0.65 on production data.
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### But...

How good are these estimates? Do we have any guarantees? Specifically, what about patient  $x_{k+1}$ ? What performance should we expect from the model for this particular instance?

# We can use PAC (probably approximately correct) theory.

Gives us valid error bounds for the model.

### But...

- Bounds are on model-level don't consider whether instance is "easy" or "hard".
- Bounds tend to be large<sup>1</sup>.

<sup>&</sup>lt;sup>1</sup>I. Nouretdinov, V. Vovk, M. Vyugin, and A. Gammerman, "Pattern recognition and density estimation under the general i.i.d. assumption," in *Computational Learning Theory*, ser. Lecture Notes in Computer Science. Springer Berlin Heidelberg, 2001, vol. 2111, pp. 337–353

### We can use Bayesian learning.

Gives us calibrated error bounds on a per-instance basis.

### But...

• Only if we know the prior probabilities<sup>2</sup>.

<sup>&</sup>lt;sup>2</sup>H. Papadopoulos, V. Vovk, and A. Gammerman, "Regression conformal prediction with nearest neighbours," *Journal of Artificial Intelligence Research*, vol. 40, no. 1, pp. 815–840, 2011

## We can use Conformal Prediction.

- Individual probabilities/error bounds per instance.
- Probabilities are well-calibrated: 80% means 80%.
- $\cdot\,$  We don't need to know the priors.
- $\cdot$  We make a single assumption exchangeability (~ i.i.d.)
- $\cdot$  We can apply it to any machine learning algorithm.
- It's rigorously proven and simple to implement!
- Developed by Vladimir Vovk, Alex Gammerman & Glenn Shafer.<sup>3</sup>

<sup>&</sup>lt;sup>3</sup>V. Vovk, A. Gammerman, and G. Shafer, *Algorithmic learning in a random world*. Springer, 2005

Conformal prediction at a glance

Assume we have

- Some distribution  $Z : X \times Y$  generating examples
- Some function  $f(z) \to \mathbb{R}$

- Apply f(z) to some, say 4, examples from Z
- Call the resulting scores  $\alpha_1, \alpha_2, \alpha_3, \alpha_4$ .
  - + For simplicity,  $\alpha_1 \leq \alpha_2 \leq \alpha_3 \leq \alpha_4$

 $\alpha_1 \qquad \alpha_2 \qquad \alpha_3 \qquad \alpha_4$ 

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20%		20%		20%		20%		20%
	$\alpha_1$		$\alpha_2$		$\alpha_3$		$\alpha_4$	

 $P[f(z) \le \alpha_3] = 0.6$  $P[f(z) \le \alpha_4] = 0.8$ 

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We know  $(x_i, y_i)$  for all examples that generated  $\alpha_1, ..., \alpha_4$ , i.e., we can obtain values for  $\alpha_1, ..., \alpha_4$ .

 20%
 20%
 20%
 20%
 20%

 0.03
 0.07
 0.11
 0.13

 $P[|y_i - h(x_i)| \le 0.11] = 0.6$  $P[|y_i - h(x_i)| \le 0.13] = 0.8$ 

## Conformal prediction: intuition

### Some intuition

For a novel example, where we know  $x_i$  but not  $y_i$ , we still know that

 $P[|y_i - h(x_i)| \le 0.11] = 0.6$  $P[|y_i - h(x_i)| \le 0.13] = 0.8$ 

and can obtain  $h(x_i)$  from our regression model, e.g.  $h(x_i) = 0.3$ .

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This is actually exactly how conformal regression works!

#### When does conformal prediction work?

We already noted a few things:

- Training data and test data belong to the same distribution (they are identically distributed)
- Choice of *f*(*z*) is irrelevant (w.r.t. validity), as long as it is symmetric (training patterns and test patterns are treated equally)

Conformal predictors output multi-valued prediction regions

 $\cdot\,$  Sets of labels or real-valued intervals

## Given

- a test pattern  $x_i$ , and
- $\cdot$  a significance level  $\epsilon$

## A conformal predictor outputs

• A prediction region  $\Gamma_i^{\epsilon}$  that contains  $y_i$  with probability  $1 - \epsilon$ 

$$Y_c = \{iris\_setosa, iris\_versicolor, iris\_virginica\}$$
  
 $Y_r = \mathbb{R}$ 

## Conformal prediction at a glance

### Point predictions

h<sub>c</sub>(x<sub>k+1</sub>) = iris\_setosa h<sub>c</sub>(x<sub>k+2</sub>) = iris\_versicolor h<sub>c</sub>(x<sub>k+3</sub>) = iris\_virginica

> $h_r(x_{k+1}) = 0.3$  $h_r(x_{k+2}) = 0.2$  $h_r(x_{k+3}) = 0.6$

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> $h_r(x_{k+1}) = 0.3$  $h_r(x_{k+2}) = 0.2$  $h_r(x_{k+3}) = 0.6$

$$P[y_i = h_c(x_i)] = ?$$
  
$$\Delta[y_i, h_r(x_i)] = ?$$

### **Prediction regions**

$$\begin{split} h_c(x_{k+1}) &= \{iris\_setosa\} \\ h_c(x_{k+2}) &= \{iris\_setosa, iris\_versicolor\} \\ h_c(x_{k+3}) &= \{iris\_setosa, iris\_versicolor, iris\_virginica\} \end{split}$$

 $h_r(x_{k+1}) = [0.2, 0.4]$  $h_r(x_{k+2}) = [0, 0.5]$  $h_r(x_{k+3}) = [0.5, 0.7]$ 

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$$\begin{split} h_c(x_{k+1}) &= \{iris\_setosa\}\\ h_c(x_{k+2}) &= \{iris\_setosa, iris\_versicolor\}\\ h_c(x_{k+3}) &= \{iris\_setosa, iris\_versicolor, iris\_virginica\} \end{split}$$

$$h_r(x_{k+1}) = [0.2, 0.4]$$
  

$$h_r(x_{k+2}) = [0, 0.5]$$
  

$$h_r(x_{k+3}) = [0.5, 0.7]$$

$$P[y_i \in h_c(x_i)] = 1 - \epsilon$$
$$P[y_i \in h_r(x_i)] = 1 - \epsilon$$

### To perform conformal prediction, we need

- A function  $f(z) \to \mathbb{R}$
- A set of training examples,  $Z^k \subset Z : X^n \times Y$
- A statistical test

## Overall rationale

- 1. Apply f(z) to training examples in  $Z^k$ , estimate distribution of  $f(z) \sim Q$
- 2. For every possible output  $\tilde{y} \in Y$ , apply f(z) to  $(x_{k+1}, \tilde{y})$
- 3. Reject  $\tilde{y}$  if it appears unlikely that  $f[(x_{k+1}, \tilde{y})] \sim Q$

# Conformal prediction at a glance

### The function *f*(*z*) We call this the nonconformity function

- A function that measures the "strangeness" of a pattern  $(x_i, y_i)$
- Any function  $f(z) \rightarrow \mathbb{R}$  works (produces valid predictions)

Properties of a good nonconformity function (that produces small prediction sets)

- Give low scores to patterns  $(x_i, y_i)$
- Give large scores to patterns  $(x_i, \neg y_i)$

Common choice:  $f(z) = \Delta[h(x_i), y_i]$ 

- *h* is called the underlying model
- "Our random forest misclassified this example, it must be weird!"

### Probability estimate for correct class

If the probability estimate for an example's correct class is low, the example is strange.

## Margin of a probability estimating model

If an example's true class is not clearly separable from other classes, it is strange.

# Distance to neighbors with same class (or distance to neighbors with different classes)

If an example is not surrounded by examples that share its label, it is strange.

## Absolute error of a regression model

If the prediction is far from the true value, the example is strange.

### rand(0, 1) Even if it's not useful, it's still valid.

### Conformal prediction process

- 1. Define a nonconformity function.
- 2. Measure the nonconformity of labeled examples  $(x_1, y_1), ..., (x_k, y_k)$ .
- 3. For a new pattern  $x_i$ , test all possible outputs  $\tilde{y} \in Y$ :
  - 3.1 Measure the nonconformity of  $(x_i, \tilde{y})$ .
  - 3.2 Is  $(x_i, \tilde{y})$  particularly nonconforming compared to the training examples? Then  $\tilde{y}$  is probably an incorrect prediction. Otherwise, include it in the prediction region.

$$p_{i}^{\tilde{y}} = \frac{\left|\left\{z_{j} \in Z : \alpha_{j} > \alpha_{i}^{\tilde{y}}\right\}\right|}{k+1} + \theta \frac{\left|\left\{z_{j} \in Z : \alpha_{j} = \alpha_{i}^{\tilde{y}}\right\}\right| + 1}{k+1}, \theta \sim U[0, 1]$$

(Portion of examples at least as nonconforming as the tentatively labeled test example)

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Prediction region

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Prediction region

$$\overline{f}_{i}^{\epsilon} = \left\{ \tilde{y} \in \mathsf{Y} : p_{i}^{\tilde{y}} > \epsilon \right\}$$

- Classification known  $\alpha_i^{\tilde{y}}$ , find  $p_i^{\tilde{y}}$
- Regression known  $p_i^{\tilde{y}}$ , find  $\alpha_i^{\tilde{y}}$

# Transductive conformal prediction (TCP) – f(z, Z)

Original conformal prediction approach

- Requires retraining model for each new test example
- $\cdot\,$  For regression problems, only certain models (e.g. kNN) can be used as of yet

# Inductive conformal prediction (ICP) - f(z)

Revised approach

- Requires model to be trained only once
- Requires that some data is set aside for calibration
  - $\cdot$  To avoid violating exchangeability assumption

Conformal classification

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**Choose an** f(z), e.g.  $f(z_i) = 1 - \hat{P}_h(y_i | x_i)$ This is the nonconformity function

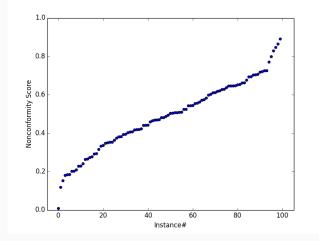
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Apply f(Z) to  $\forall z_i \in Z_c$ Save these calibration scores We denote these  $\alpha_1, ..., \alpha_q$ 

Apply f(z) to  $Z_c$ , and obtain a set of calibration scores  $\alpha_1, ..., \alpha_q$ 



For each  $\tilde{y} \in Y$ Let  $\alpha_i^{\tilde{y}} = f[(x_i, \tilde{y})]$ 

Calculate

$$p_{i}^{\tilde{y}} = \frac{\left|\left\{z_{j} \in Z_{c} : \alpha_{j} > \alpha_{i}^{\tilde{y}}\right\}\right|}{q+1} + \theta \frac{\left|\left\{z_{j} \in Z_{c} : \alpha_{j} = \alpha_{i}^{\tilde{y}}\right\}\right| + 1}{q+1}, \theta \sim U[0, 1]$$

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Fix a significance level  $\epsilon \in (0, 1)$ 

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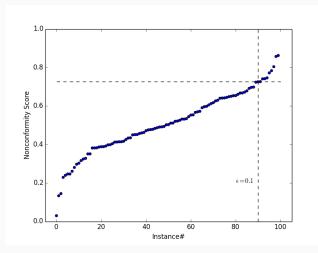
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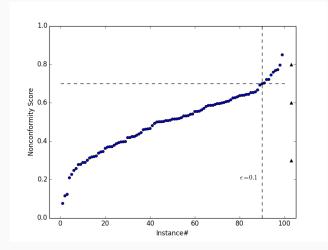
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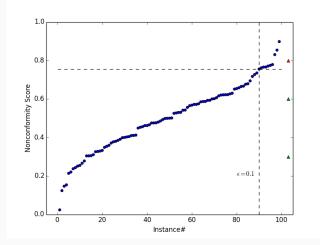
### Choose a significance level $\epsilon$



Obtain  $\alpha_i$  using f(z) for each possible class  $(x_i, \tilde{y}_1), (x_i, \tilde{y}_2), (x_1, \tilde{y}_3), ...,$  resulting in  $\alpha_i^{\tilde{y}_1}, \alpha_i^{\tilde{y}_2}, \alpha_i^{\tilde{y}_3}, ...$ 



Reject/include based on the *p*-value statistic, and the chosen  $\epsilon$ 



### Predicting whether a customer will churn or not - a real-world example

- A data set from one of the leading e-retailers in Sweden consisting of altogether 255298 customers.
- The target variable for the analysis is whether the specific customer will churn or not, i.e., no purchase one year after the previous order.
- Each customer is described using altogether 276 attributes.
- We are not allowed to give a detailed description of all the attributes, but they include statistics like number of orders, number of visits to the website and whether the customer has clicked on promotion emails sent by the retailer.

### Predicting whether a customer will churn or not - 16 sample instances

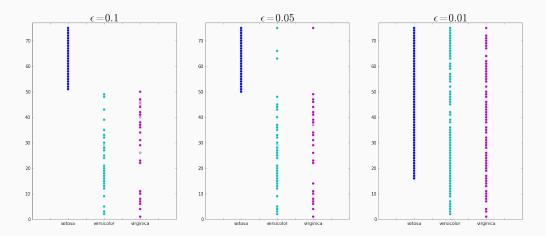
Correct	$\epsilon = 0.2$	$\epsilon = 0.1$	$\epsilon = 0.05$	$\epsilon = 0.01$
Churn	{Churn}	{Churn}	{Churn}	{Churn}
Loyal	{Churn}	{Churn}	{Loyal, Churn}	{Loyal, Churn}
Loyal	{}	{Loyal}	{Loyal}	{Loyal}
Churn	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}
Churn	{Churn}	{Churn}	{Loyal, Churn}	{Loyal, Churn}
Churn	{Churn}	{Churn}	{Churn}	{Loyal, Churn}
Loyal	{Loyal}	{Loyal}	{Loyal, Churn}	{Loyal, Churn}
Churn	{Churn}	{Churn}	{Churn}	{Churn}
Loyal	{Loyal}	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}
Loyal	{Loyal}	{Loyal}	{Loyal}	{Loyal, Churn}
Churn	{Churn}	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}
Churn	{Churn}	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}
Loyal	{Loyal}	{Loyal}	{Loyal}	{Loyal}
Churn	{Loyal}	{Loyal}	{Loyal}	{Loyal, Churn}
Loyal	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}	{Loyal, Churn}
Loyal	{Loyal}	{Loyal}	{Loyal}	{Loyal, Churn}

#### Predicting whether a customer will churn or not - overall results

	$\epsilon = 0.2$	$\epsilon = 0.1$	$\epsilon = 0.05$	$\epsilon = 0.01$
RF 300				
AvgC	1.061	1.334	1.519	1.791
OneC	0.939	0.666	0.481	0.209
Errors	0.202	0.100	0.052	0.010
LogReg				
AvgC	1.075	1.347	1.525	1.790
OneC	0.925	0.653	0.475	0.210
Errors	0.199	0.096	0.050	0.011

- For classification, an error is when the correct label is not in the prediction set, i.e., for two-class problems incorrect singleton predictions and empty predictions.
- The probability for an error is always the chosen  $\epsilon$ .
- $\cdot$  An obvious and user-controlled trade-off between errors and prediction size

### Iris, Random Forest



Conformal regression

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## Divide the training set Z into two disjoint subsets

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Apply f(z) to  $\forall z_i \in Z_c$ Save these calibration scores, sorted in descending order We denote these  $\alpha_1, ..., \alpha_q$ 

## Inductive Conformal Regression

Fix a significance level  $\epsilon \in (0, 1)$ 

Let  $s = \lfloor \epsilon(q+1) \rfloor$ .

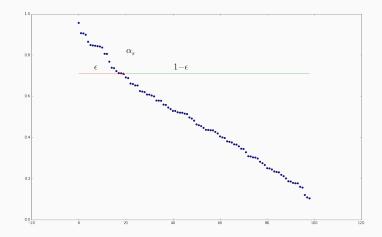
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The interval contains  $y_i$  with probability  $1 - \epsilon$ 

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#### Note

For regression, we can't enumerate each  $\tilde{y} \in Y$ , instead we work backwards, i.e., fix the *p*-value and then find an appropriate  $\alpha_i^{\tilde{y}}$ .

• Hence, our nonconformity function must be (partially) invertible for quick calculation of intervals

A sample regression problem - Boston Housing Attributes:

- CRIM: per capita crime rate by town
- ZN: proportion of residential land zoned for lots over 25,000 sq.ft.
- INDUS: proportion of non-retail business acres per town
- CHAS: Charles River dummy variable (= 1 if tract bounds river; 0 otherwise)
- NOX: nitric oxides concentration (parts per 10 million)
- RM: average number of rooms per dwelling
- AGE: proportion of owner-occupied units built prior to 1940
- DIS: weighted distances to five Boston employment centres
- RAD: index of accessibility to radial highways
- TAX: full-value property-tax rate per \$10000
- PTRATIO: pupil-teacher ratio by town
- B:  $1000(Bk 0.63)^2$  where Bk is the proportion of blacks by town
- LSTAT: % lower status of the population
- Price

# Inductive Conformal Regression

## Predicting price - 16 sample instances

	$\epsilon =$	0.2	$\epsilon = 0.1$		$\epsilon = 0.05$		$\epsilon = 0.01$	
Correct	Min	Max	Min	Max	Min	Max	Min	Max
10.8	6.7	23.2	2.7	27.3	0.0	31.0	0.0	40.7
14.9	9.9	26.4	5.8	30.4	2.1	34.1	0.0	43.8
12.6	10.4	26.3	6.6	30.1	3.0	33.7	0.0	43.0
14.9	16.8	30.2	13.5	33.5	10.5	36.5	2.6	44.4
19.1	9.2	25.6	5.2	29.6	1.5	33.3	0.0	43.0
20.1	11.7	28.1	7.7	32.1	4.1	35.8	0.0	45.4
19.9	10.2	26.5	6.2	30.5	2.5	34.2	0.0	43.9
23	12.9	29.2	8.9	33.2	5.2	36.9	0.0	46.6
23.7	20.5	36.4	16.7	40.2	13.1	43.8	3.8	53.1
21.8	13.1	28.5	9.4	32.2	6.0	35.7	0.0	44.7
20.6	13.0	29.4	9.0	33.4	5.3	37.1	0.0	46.7
19.1	11.1	27.4	7.1	31.4	3.4	35.1	0.0	44.8
15.2	10.3	26.8	6.3	30.8	2.6	34.5	0.0	44.3
7.0	7.7	24.2	3.6	28.2	0.0	31.9	0.0	41.6
24.5	18.0	23.4	16.6	24.8	15.4	26.0	12.2	29.2
11.9	17.8	24.1	16.3	25.6	14.9	27.1	11.1	30.8

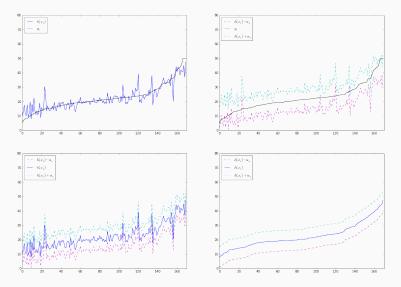
### Overall results

	$\epsilon = 0.2$	$\epsilon = 0.1$	$\epsilon = 0.05$	$\epsilon = 0.01$
Errors	0.201	0.090	0.053	0.011
Average interval	10.1	16.0	19.4	32.8

- For regression problems, an error is when the target variable is outside of the interval.
- The probability for an error is always the chosen  $\epsilon$ .
- Again an obvious and user-controlled trade-off between errors and prediction size
- $\cdot$  This data set is rather small, so the empirical error rates differ slightly from  $\epsilon$

# Inductive Conformal Regression

Boston Housing, Random Forest,  $\epsilon = 0.1$ 



### Static prediction interval size

Using  $f(z_i) = |y_i - h(x_i)|$  and  $\Gamma_i^{\epsilon} = h(x_i) \pm \alpha_s$ means each prediction interval has the same size  $(\alpha_s)$ .

But we want individual bounds for each x<sub>i</sub>...

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### Normalized nonconformity functions

Normalized nonconformity functions utilize an additional term  $\sigma_i$ .

$$f(z_i) = \frac{|y_i - h(x_i)|}{\sigma_i}$$

 $\sigma_i$  is an estimate of the difficulty of predicting  $y_i$ 

A common practice is to let  $\sigma$  be predicted by a model, e.g.,  $\sigma_i = \hat{\Delta}[y_i, h(x_i)]$ 

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The normalized prediction for a new example is  $\Gamma_i^{\epsilon} = h(x_i) \pm \alpha_s \sigma_i$ 

Divide the training set Z into two disjoint subsets A proper training set  $Z_t$ A calibration set  $Z_c$ 

Fit a model h using  $Z_t$ In addition

- Let  $E_t$  be the residual errors of h (i.e. the errors that h makes on  $Z_t$ )
- Fit a model g using  $X_t \times E_t$

$$f(z_i) = \frac{|y_i - h(x_i)|}{g(x_i) + \beta}$$

 $\boldsymbol{\beta}$  is a sensitivity parameter that determines the impact of normalization

Apply f(z) to  $\forall z_i \in Z_c$ Save these calibration scores, sorted in descending order

## Fix a significance level $\epsilon \in (0, 1)$

Let  $s = \lfloor \epsilon(q+1) \rfloor$ This is the index of the  $(1 - \epsilon)$ -percentile nonconformity score,  $\alpha_s$ .

## Prediction region

The prediction for a new example is  $\Gamma_i^{\epsilon} = h(x_i) \pm \alpha_s(g(x_i) + \beta)$ Interval contains  $y_i$  with probability  $1 - \epsilon$ 

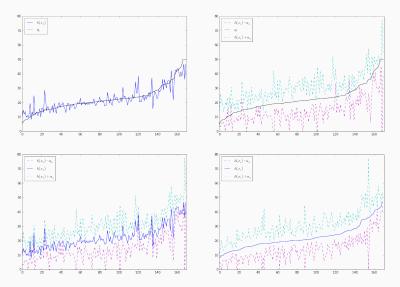
## Effects of normalization

Normalization produces more specific (individualized) predictions.

The intervals tend to be tighter, on average, when using normalization.

# Inductive Conformal Regression

Boston Housing, Random Forest, normalized nonconformity function,  $\epsilon=$  0.1



Validity and efficiency

### Conformal predictors are subject to two desiderata

Validity — coherence between  $\epsilon$  and error rate Efficiency — size of prediction regions (i.e. informativeness)

Conformal predictors are automatically valid

Efficiency depends on the nonconformity function (and thus the underlying model)

### Conformal predictors are subject to two desiderata

Validity — coherence between  $\epsilon$  and error rate Efficiency — size of prediction regions (i.e. informativeness)

Conformal predictors are automatically valid Efficiency depends on the nonconformity function (and thus the underlying model)

## Confidence-efficiency trade-off

The more confidence we require in a prediction, the larger the prediction region will be

$\epsilon$	errors	size
0.01	0.006	38.31
0.05	0.040	16.90
0.10	0.089	11.46
0.20	0.191	7.562

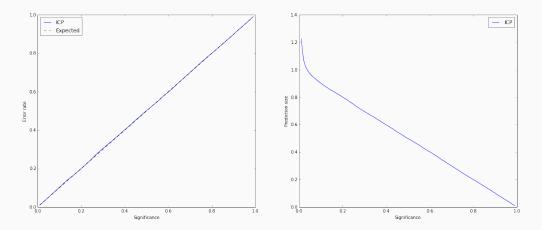
Table 1: Boston 10x10 RF CV

$\epsilon$	errors	size
0.01	0.011	2.347
0.05	0.055	1.052
0.10	0.100	0.930
0.20	0.202	0.804

Table 2: Iris 10x10 RF CV

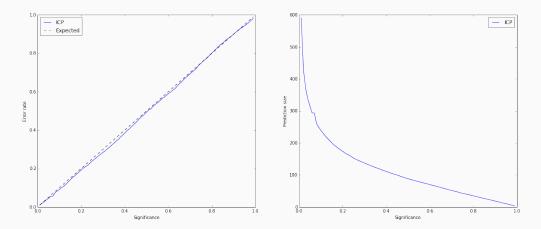
# Validity and efficiency

## Digits (classification), Random Forest, 10x10 CV



# Validity and efficiency

## Diabetes (regression), Random Forest, 10x10 CV



Empirical validity is measured by observing the error rate of a conformal predictor.

Efficiency can be measured in many different ways<sup>4</sup>.

Examples — regression

 $\cdot$  Average size of prediction interval

Examples — classification

- Average number of classes per prediction (AvgC)
- Rate of predictions containing a single class (OneC)
- Average *p*-value

<sup>&</sup>lt;sup>4</sup>V. Vovk, V. Fedorova, I. Nouretdinov, and A. Gammerman, "Criteria of efficiency for conformal prediction," 2014

Considerations and modifications

## Conformal predictors are, by default, unconditional

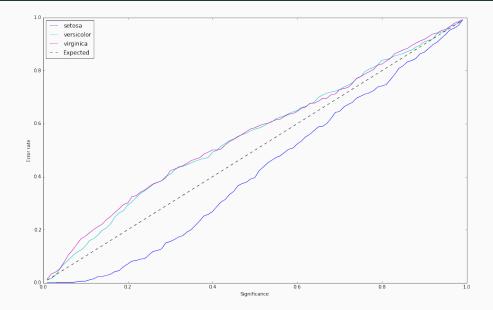
Their guaranteed error rate applies to the entire test set.

- $\cdot\,$  Difficult patterns (e.g. minority class) may see a greater error rate than expected
- Easy patterns (e.g. majority class) may see a smaller error rate than expected

## Example — Iris data set

- $\cdot$  One linearly separable class (easy)
- Two linearly non-separable classes (difficult)

# Conditional conformal prediction



## Conditional conformal predictors<sup>5</sup> help solve this by

Dividing the problem space into several disjoint subspaces

- $\cdot\,$  e.g. let each class represent a subspace, or
- define subspace based on some input variable(s) (age, gender, etc.)

Guaranteeing an error rate at most  $\epsilon$  for each subspace

<sup>&</sup>lt;sup>5</sup>V. Vovk, "Conditional validity of inductive conformal predictors," *Journal of Machine Learning Research -Proceedings Track*, vol. 25, pp. 475–490, 2012

# Conditional conformal prediction

# **Define a mapping function** $K(z_i) = \kappa_i$ Examples

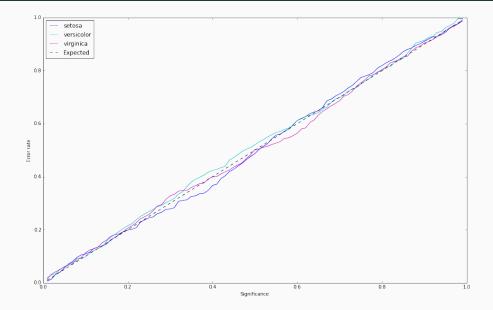
$$K(z_i) = y_i \tag{1}$$

$$K(z_i) = \begin{cases} 1 & \text{if } x_{i,1} < 50 \\ 2 & \text{if } 50 \le x_{i,1} < 100 \\ 3 & \text{otherwise} \end{cases}$$
(2)

#### Conditional *p*-value

$$p_{i}^{\tilde{y}} = \frac{|\{z_{j} \in Z_{c} : \alpha_{j} > \alpha_{i}^{\tilde{y}}\} \land K(z_{i}) = K(z_{j})|}{|K(z_{i}) = K(z_{j})| + 1} + \theta \frac{|\{z_{j} \in Z_{c} : \alpha_{j} = \alpha_{i}^{\tilde{y}}\} \land K(z_{i}) = K(z_{j})|}{|K(z_{i}) = K(z_{j})| + 1}, \theta \sim U[0, 1]$$

# Conditional conformal prediction



### The calibration set

Inductive conformal predictors need some data set aside for calibration? — How much?

25%  $\sim$  33% are common choices, and provide a good balance between underlying model performance and calibration accuracy<sup>6</sup>.

### Alternatives

Bagged ensembles can use out-of-bag examples for calibration<sup>7 8</sup>.

<sup>&</sup>lt;sup>6</sup>H. Linusson, U. Johansson, H. Boström, and T. Löfström, "Efficiency comparison of unstable transductive and inductive conformal classifiers," in *Artificial Intelligence Applications and Innovations*. Springer, 2014, pp. 261–270

<sup>&</sup>lt;sup>7</sup>U. Johansson, H. Boström, T. Löfström, and H. Linusson, "Regression conformal prediction with random forests," *Machine Learning*, vol. 97, no. 1-2, pp. 155–176, 2014

<sup>&</sup>lt;sup>8</sup>H. Boström, H. Linusson, T. Löfström, and U. Johansson, "Accelerating difficulty estimation for conformal regression forests," *Annals of Mathematics and Artificial Intelligence*, pp. 1–20, 2017

## The calibration set cont.

For an inductive conformal predictor to be exactly valid, it requires exactly  $k\epsilon^{-1} - 1$  calibration instances.

- Otherwise, discretization errors come into play
  - (Rendering the conformal predictor conservatively valid)
- $\cdot$  Of particular importance when calibration set is small
  - $\cdot\,$  e.g. when using conditional conformal prediction

## Alternatives

Interpolation of *p*-values can alleviate this problem.<sup>9</sup> <sup>10</sup>

<sup>&</sup>lt;sup>9</sup>L. Carlsson, E. Ahlberg, H. Boström, U. Johansson, and H. Linusson, "Modifications to p-values of conformal predictors," in *Statistical Learning and Data Sciences*. Springer, 2015, pp. 251–259

<sup>&</sup>lt;sup>10</sup>U. Johansson, E. Ahlberg, H. Boström, L. Carlsson, H. Linusson, and C. Sönströd, "Handling small calibration sets in mondrian inductive conformal regressors," in *Statistical Learning and Data Sciences*. Springer, 2015, pp. 271–280

Conformal classification - a critical look

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- $\cdot\,$  Thus, all errors must be made on the remaining singleton predictions.
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- Thus, all errors must be made on the remaining singleton predictions.
- So, once we have observed a singleton prediction, the probability for that being incorrect is most likely much higher than  $\epsilon$ .
- $\cdot\,$  It must be noted that this "problem" does not exist in conformal regression.

# Venn predictors

#### Introduction

- Many classifiers are able to output not only the predicted class label, but also a probability distribution over the possible classes.
- Naturally, all probabilistic prediction requires that the probability estimates are well-calibrated, i.e., the predicted class probabilities must reflect the true, underlying probabilities.
- If this is not the case, the probabilistic predictions actually become misleading.

#### Calibration

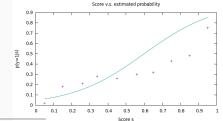
- In probabilistic prediction, the task is to predict the probability distribution of the label, given the training set and the test object.
- The goal is to obtain a valid predictor.
- In general, validity means that the probability distributions from the predictor must perform well against statistical tests based on subsequent observation of the labels.
- We are interested in calibration:  $p(c_j | p^{c_j}) = p^{c_j}$ , where  $p^{c_j}$  is the probability estimate for class *j*.

## Platt scaling

Platt scaling<sup>11</sup> was originally introduced as a method for calibrating support-vector machines. It works by finding the parameters of a sigmoid function maximizing the likelihood of a calibration set. The function is

$$\hat{p}(c \mid s) = \frac{1}{1 + e^{As + B}},$$
(3)

where  $\hat{p}(c \mid s)$  gives the probability that an example belongs to class *c*, given that it has obtained the score *s*, and where *A* and *B* are parameters of the function found by gradient descent search.



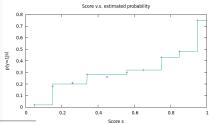
<sup>11</sup>J. C. Platt, "Probabilistic outputs for support vector machines and comparisons to regularized likelihood methods," in *Advances in Large Margin Classifiers*. MIT Press, 1999, pp. 61–74

### Isotonic regression

Isotonic regression<sup>12</sup> is a calibration method that can be regarded as a general form of binning, not requiring a predetermined number of bins.

The calibration function, which is assumed to be isotonic, i.e., non-decreasing, is a step-wise regression function, which can be learned by an algorithm known as the pair-adjacent violators algorithm.

The algorithm outputs a function that for each input probability interval returns the fraction of positive examples in the calibration set in that interval.



<sup>12</sup>B. Zadrozny and C. Elkan, "Obtaining calibrated probability estimates from decision trees and naive Bayesian classifiers," in *Proc. 18th International Conference on Machine Learning*, 2001, pp. 609–616

Venn predictors<sup>13</sup>, are multi-probabilistic predictors with proven validity properties.

Venn predictors was originally suggested in a transductive setting, but here we present the inductive variant:

To construct an inductive Venn predictor, the available labeled training examples  $(\{(x_1, y_1), \ldots, (x_l, y_l)\})$  are split into two parts, the proper training set  $(\{(x_1, y_1), \ldots, (x_q, y_q)\})$ , used to train an underlying model, and a calibration set  $(\{(x_{q+1}, y_{q+1}), \ldots, (x_l, y_l)\})$  used to estimate label probabilities for each new test example.

When presented with a new test object  $x_{l+1}$ , the aim of Venn prediction is to estimate the probability that  $y_{l+1} = Y_i$ , for each  $Y_i$  in the set of possible labels  $Y_i \in \{Y_1, \dots, Y_c\}$ .

<sup>&</sup>lt;sup>13</sup>V. Vovk, G. Shafer, and I. Nouretdinov, "Self-calibrating probability forecasting," in Advances in Neural Information Processing Systems, 2004, pp. 1133–1140

The key idea of Venn prediction is to divide all calibration examples into a number of k categories and use the relative frequency of label  $Y_j \in \{Y_1, \ldots, Y_c\}$  in each category to estimate label probabilities for test instances falling into that category.

The categories are defined using a Venn taxonomy and every taxonomy leads to a different Venn predictor.

Typically, the taxonomy is based on the underlying model, trained on the proper training set, and for each calibration and test object  $x_i$ , the output of this model is used to assign  $(x_i, y_i)$  into one of the categories.

One basic Venn taxonomy, which can be used with every kind of classification model, simply puts all examples predicted with the same label into the same category.

For test instances, the category is first determined using the underlying model, in an identical way as for the calibration instances. Then, the label frequencies of the calibration instances in that category are used to calculate the estimated label probabilities.

As in conformal prediction, the test instance  $z_{l+1}$  is included in this calculation. However, since the true label  $y_{l+1}$  is not known for the test object  $x_{l+1}$ , all possible labels  $Y_i \in \{Y_1, \ldots, Y_c\}$  are used to create a set of label probability distributions.

Instead of dealing directly with these distributions, the lower  $L(Y_j)$  and upper  $U(Y_j)$  probability estimates for each label  $Y_j$  are often used.

Let *k* be the category assigned to the test object  $x_{l+1}$  by the Venn taxonomy, and  $Z_k$  be the set of calibration instances belonging to category *k*. Then the lower and upper probability estimates are defined by:

$$L(Y_j) = \frac{|\{(x_m, y_m) \in Z_k \mid y_m = Y_j\}|}{|Z_k| + 1}$$
(4)

and:

$$U(Y_j) = \frac{\left|\{(x_m, y_m) \in Z_k \mid y_m = Y_j\}\right| + 1}{|Z_k| + 1}$$
(5)

In order to make a prediction  $\hat{y}_{l+1}$  for  $x_{l+1}$  using the lower and upper probability estimates, the following procedure is often employed:

$$\hat{y}_{l+1} = \max_{Y_j \in \{Y_1, \dots, Y_c\}} L(Y_j)$$
(6)

The output of a Venn predictor is the above prediction  $\hat{y}_{l+1}$  together with the probability interval:

$$[L(\hat{y}_{l+1}), U(\hat{y}_{l+1})] \tag{7}$$

Nonconformist - conformal prediction in Python

#### How good is your prediction?

You want to estimate the risk of cancer recurrence in patient  $x_{k+1}$ 

To your disposal, you have:

- 1. A set of historical observations  $(x_1, y_1), \ldots, (x_k, y_k)$ 
  - x<sub>i</sub> describes a patient by age, tumor size, etc
  - $y_i$  is a measurement of cancer recurrence in patient  $x_i$
- 2. Some machine learning (classification or regression) algorithm
- 3. Conformal prediction

```
import pandas as pd
```

```
breast_cancer = pd.read_csv('./data/breast-cancer.csv')
```

```
# proper training set
x_train = breast_cancer.values[:-100, :-1]
y_train = breast_cancer.values[:-100, -1]
```

```
# calibration set
x_cal = breast_cancer.values[-100:-1, :-1]
y_cal = breast_cancer.values[-100:-1, -1]
```

```
# (x_k+1, y_k+1)
x_test = breast_cancer.values[-1, :-1]
y_test = breast_cancer.values[-1, -1]
```

# Omitted: convert y\_train, y\_cal, y\_test to numeric

```
import numpy as np
from sklearn.neighbors import KNeighborsClassifier
from nonconformist.icp import IcpClassifier
from nonconformist.nc import NcFactory
```

```
knn = KNeighborsClassifier(n_neighbors=5)
nc = NcFactory.create_nc(knn)
icp = IcpClassifier(nc)
```

```
icp.fit(x_train, y_train)
icp.calibrate(x_cal, y_cal)
```

print(icp.predict(np.array([x\_test]), significance=0.05))

[[ True False ]]

## Nonconformist

Installation options:

- git clone http://github.com/donlnz/nonconformist
- pip install nonconformist

Nonconformist supports:

- Conformal classification (inductive)
- Conformal regression (inductive)
- Mondrian (e.g., class-conditional) models
- Normalization
- · Aggregated conformal predictors ( $\approx$  icp ensembles)
- Out-of-bag calibration
- Plug-and-play using sklearn
- User extensions

### Questions, suggestions, feedback, contributions, etc.?

henrik.linusson@hb.se

Other scenarios and suggested reading

#### Other scenarios for conformal prediction

- Anomaly detection with guaranteed maximum false positive rates.<sup>14</sup>
- Concept drift detection / i.i.d. checking with maximum false positive rates.<sup>15</sup>
- Rule exctraction with guaranteed fidelity.<sup>16</sup>
- Semi-supervised learning.<sup>17</sup>

<sup>&</sup>lt;sup>14</sup>R. Laxhammar and G. Falkman, "Conformal prediction for distribution-independent anomaly detection in streaming vessel data," in *Proceedings of the First International Workshop on Novel Data Stream Pattern Mining Techniques*. ACM, 2010, pp. 47–55

<sup>&</sup>lt;sup>15</sup>V. Fedorova, A. Gammerman, I. Nouretdinov, and V. Vovk, "Plug-in martingales for testing exchangeability on-line," in *29th International Conference on Machine Learning*, 2012

<sup>&</sup>lt;sup>16</sup>U. Johansson, R. König, H. Linusson, T. Löfström, and H. Boström, "Rule extraction with guaranteed fidelity," in *Artificial Intelligence Applications and Innovations*. Springer, 2014, pp. 281–290

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#### Not (yet) proven valid

But seems to be working well in practice.

#### Application domains

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- I. Nouretdinov, S. G. Costafreda, A. Gammerman, A. Chervonenkis, V. Vovk, V. Vapnik, and C. H. Fu, "Machine learning classification with confidence: application of transductive conformal predictors to mri-based diagnostic and prognostic markers in depression," *Neuroimage*, vol. 56, no. 2, pp. 809–813, 2011
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## **Questions?**

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