Week 13: Conformal Prediction MATH-517 Statistical Computation and Visualization

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Motivation

- Prediction algorithms usually return point predictions
- In practice we also need a notion of uncertainty:
 - prediction intervals,
 - or more general prediction sets
- Classical approaches:
 - parametric models + asymptotic normality
 - bootstrap, jacknife, Bayesian intervals, etc
- Today: conformal prediction
 - wraps around any black-box predictor
 - gives finite-sample, distribution-free guarantees

Abmbitious aim

We observe i.i.d. training data $(X_i,Y_i)\sim P$, $i=1,\dots,n$

Given a miscoverage level $\alpha \in [0,1]$, we want a set-valued predictor

$$\hat{C}_\alpha:\mathcal{X}\to\{\text{subsets of }\mathbb{R}\}$$

such that, for a new i.i.d. pair (X_{n+1},Y_{n+1}) ,

$$\mathbb{P}\big(Y_{n+1} \in \hat{C}_{\alpha}(X_{n+1})\big) \geq 1 - \alpha$$

for any distribution P

- No asymptotics, i.e., finite sample guarantees
- ullet No parametric assumptions (agnostic to P and the model/algorithm)
- Sets should still be reasonably short when prediction is easy

Exchangeability

• A sequence of random variables (Z_1,Z_2,\ldots,Z_{n+1}) is **exchangeable** if its joint distribution does not change when we permute the indices, i.e., for any permutation π :

$$(Z_1,\ldots,Z_{n+1})\stackrel{d}{=}(Z_{\pi(1)},\ldots,Z_{\pi(n+1)})$$

- i.i.d. samples are always exchangeable, but exchangeability is slightly more general (think of $\mathcal{N}_d(\mu,\Sigma)$, with $\sigma_{ij}=\delta^2$, for $i\neq j$)
- Symmetry under permutations ⇒ certain order-statistic arguments (e.g., about ranks or quantiles) hold exactly in finite samples: this is the key ingredient behind conformal prediction

Consequence of exchangeability I (Vovk et al., 2005)

Start with exchangeable Y_1,\dots,Y_n , and let $Y_{(1)} \leq \dots \leq Y_{(n)}$ denote their order statistics

For a target miscoverage level $\alpha \in (0,1)$, define the empirical $(1-\alpha)$ -quantile as

$$\hat{q}_{1-\alpha} = \begin{cases} Y_{(\lceil (n+1)(1-\alpha)\rceil)}, & \text{if } \lceil (1-\alpha)(n+1)\rceil \leq n, \\ \infty, & \text{otherwise} \end{cases}$$

For one more exchangeable sample Y_{n+1} , we obtain

$$\mathbb{P}\big(Y_{n+1} \le \hat{q}_{1-\alpha}\big) \ \ge \ 1 - \alpha$$

Additionally, under no ties, i.e., Y_1,\ldots,Y_n,Y_{n+1} are a.s. distinct,

$$\mathbb{P}(Y_{n+1} \leq \hat{q}_{1-\alpha}) < (1-\alpha) + \frac{1}{n+1}$$

Consequence of exchangeability II

 \bullet Exchangeability \Rightarrow the rank of Y_{n+1} among (Y_1,\ldots,Y_n,Y_{n+1}) is uniformly distributed over $\{1,\ldots,n+1\}.$ Thus

$$\begin{split} \mathbb{P}(\mathrm{Rank}_{n+1} \leq \lceil (1-\alpha)(n+1) \rceil) &= \sum_{j=1}^{\lceil (1-\alpha)(n+1) \rceil} \mathbb{P}(\mathrm{Rank}_{n+1} = j) \\ &\geq \frac{\lceil (1-\alpha)(n+1) \rceil}{n+1} \, \geq \, 1-\alpha \\ &\text{a.s. } \underset{=}{\mathrm{distinct}} \frac{\lceil (1-\alpha)(n+1) \rceil}{n+1} < \frac{(1-\alpha)(n+1) + 1}{n+1} \end{split}$$

• Note that $\hat{q}_{1-\alpha}$ is computed over Y_1,\dots,Y_n thanks to the equivalence between

$$\mathbb{P}(Y_{n+1} \text{ is among the } \lceil (1-\alpha)(n+1) \rceil \text{ smallest of } Y_1,\dots,Y_{n+1}) \geq 1-\alpha$$

$$\mathbb{P}(Y_{n+1} \text{ is among the } \lceil (1-\alpha)(n+1) \rceil \text{ smallest of } Y_1,\dots,Y_n) \geq 1-\alpha$$

Section 1

Naive attempt for regression

Naive construction

• Fit any regression algorithm on the entire (training) data set:

$$\hat{f}_n:\mathcal{X}\to\mathbb{R}$$

Training residuals:

$$R_i = |Y_i - \hat{f}_n(X_i)|, \quad i = 1, \dots, n$$

- \bullet Let $\hat{q}_{1-\alpha}$ be the empirical $(1-\alpha)\text{-quantile}$ of $\{R_i\}$
- For a test point $X_{n+1} = x$, define prediction set

$$\hat{C}_{\alpha}(x) = \{y: \mid y - \hat{f}_n(x) \mid \leq \hat{q}_{1-\alpha}\} = [\,\hat{f}_n(x) - \hat{q}_{1-\alpha}, \,\, \hat{f}_n(x) + \hat{q}_{1-\alpha}\,]$$

Looks reasonable, but:

the test residual

$$R_{n+1} = |Y_{n+1} - \hat{f}_n(X_{n+1})|$$

is not exchangeable with R_1,\dots,R_n , because \hat{f}_n was trained on (X_i,Y_i) but not on (X_{n+1},Y_{n+1})

ullet typically we get under-coverage (especially if \hat{f}_n overfits the training data)

Recover exchangeability

To recover exchangeability we must:

- construct conformity scores (residuals) that treat calibration points and the test point symmetrically
- rely on data splitting or cross-validation

This leads to split conformal prediction (SCP)

Section 2

Split Conformal Prediction (Regression)

Algorithm

Split index set into two disjoint parts:

- proper training set D_1
- calibration set D₂
- Fit a point predictor \hat{f}_{n_i} using only $(X_i, Y_i)_{i \in D_i}$
- 2 For $i \in D_2$, define the conformity scores (calibration residuals)

$$R_i = |Y_i - \hat{f}_{n_1}(X_i)|$$

- **3** Let $\hat{q}_{1-\alpha}$ be the $\lceil (1-\alpha)(n_2+1) \rceil$ -th smallest residual
- For a test point x (exchangeable with D_2), define the prediction interval

$$\hat{C}_{\alpha}(x) = [\,\hat{f}_{n_1}(x) - \hat{q}_{1-\alpha},\,\,\hat{f}_{n_1}(x) + \hat{q}_{1-\alpha}\,]$$

Coverage guarantee (upper bound under assumption of no ties):

$$1-\alpha \leq \mathbb{P}\big(Y_{n+1} \in \hat{C}_{\alpha}(X_{n+1})\big) \leq 1-\alpha + \frac{1}{n_2+1}$$

Marginal versus conditional coverage

Conformal prediction guarantees marginal coverage:

$$\mathbb{P}\{Y_{n+1} \in \widehat{C}_{\alpha}(X_{n+1})\} \ge 1 - \alpha,$$

averaged over the randomness in the calibration and test point (X_{n+1}, Y_{n+1})

Conditioning on the calibration set, the distribution of the coverage is

$$\mathbb{P}\left(Y_{n+1} \in \widehat{C}_{\alpha}(X_{n+1}) \mid \left\{(X_i, Y_i)\right\}_{i=1}^{n_2}\right) \sim \mathrm{Beta}(n_2 + 1 - l, l),$$

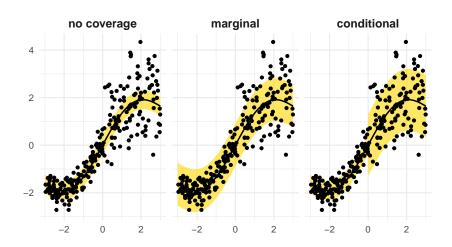
with
$$l=\lfloor (n_2+1)\alpha\rfloor \Rightarrow$$
 has mean= $\frac{\lceil (1-\alpha)(n_2+1)\rceil}{n_2+1}$ (sanity check)

This is different from conditional coverage:

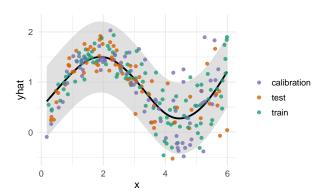
$$\Pr\{Y_{n+1} \in \widehat{C}_{\alpha}(x) \mid X_{n+1} = x\} \geq 1 - \alpha \quad \text{for all } x,$$

which is generally impossible without assumptions on the distribution of the data; see Vovk (2012) and Lei and Wasserman (2013)

Marginal versus conditional coverage



Coverage is not adaptive



- better prediction algorithms (from proper training set) lead to smaller prediction sets (width measured on average over x), but
- ullet split conformal bands with absolute residuals have width exactly constant in $x \Rightarrow$ no adaptivity to local hardness of the prediction

Section 3

Conformalised Quantile Regression (CQR)

Motivation: quantile models to improve local adaptivity

- Split conformal regression uses absolute residuals, producing constant-width bands
- ullet But often the noise varies with x: heteroskedasticity, skewness, outliers ...
- Idea: instead of modelling mean+variance, model conditional quantiles:

$$f_{\tau}(x) \approx Q_{Y|X=x}(\tau)$$

- For a prediction point x, fit:
 - ullet a lower model $f_{lpha/2}(x)$
 - an upper model $\hat{f}_{1-lpha/2}(x)$

These already form a prediction band, but with no coverage guarantee!

CQR adds calibration to guarantee finite-sample coverage

Quantile regression

For a random variable Z with cdf F, the au-quantile is

$$Q_Z(\tau) = \inf\{z : F(z) \ge \tau\}$$

In regression, we model conditional quantiles:

$$f_{\tau}^{\star}(x) = Q_{Y|X=x}(\tau)$$

Quantile regression estimates $f_{ au}^{\star}$ by minimising the pinball loss

$$\ell_\tau(y,u) = (\tau - \mathbf{1}\{y < u\})(y-u)$$

Fitting conditional quantiles on training data

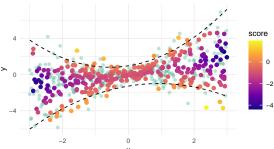
Step 1: lower and upper quantiles on training data no finite–sample guarantees

Calibration: checking how well the quantile band fits unseen data

We now compute how much calibration points fall outside the quantile band Conformity score (negatively oriented) for each point $i \in D_2$:

$$R_i = \max\{\hat{f}_{\alpha/2}(X_i) - Y_i, \ Y_i - \hat{f}_{1-\alpha/2}(X_i)\}$$

Step 2: Conformity scores on calibration data



CQR interval: quantile band expanded just enough for finite-sample validity

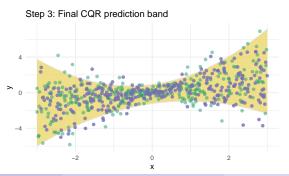
Compute the quantile of the conformity scores (on calibrated data)

$$\hat{q}_{1-\alpha} = R_{(\lceil (1-\alpha)(n_2+1)\rceil)}$$

Final interval

$$\hat{C}_{\alpha}(x) = [\hat{f}_{\alpha/2}(x) - \hat{q}_{1-\alpha}, \quad \hat{f}_{1-\alpha/2}(x) + \hat{q}_{1-\alpha}]$$

has valid finite-sample coverage, though coverage is only marginal!



CQR: Algorithm

- f 0 On the train set D_1 , fit two quantile models:
 - $\bullet \ \hat{f}_{\alpha/2}(x) \ \text{and} \ \hat{f}_{1-\alpha/2}(x) \\$
- $oldsymbol{0}$ On calibration set D_2 , define scores

$$R_i = \max\{\hat{f}_{\alpha/2}(X_i) - Y_i, Y_i - \hat{f}_{1-\alpha/2}(X_i)\}$$

- $\ \, \mathbf{0} \ \, \mathrm{Let} \,\, \hat{q}_{1-\alpha} \,\, \mathrm{be} \,\, \mathrm{the} \,\, \lceil (1-\alpha)(n_2+1) \rceil \text{-th smallest score}$
- Final prediction band:

$$\hat{C}_{\alpha}(x) = [\,\hat{f}_{\alpha/2}(x) - \hat{q}_{1-\alpha},\,\,\hat{f}_{1-\alpha/2}(x) + \hat{q}_{1-\alpha}\,], \label{eq:constraint}$$

with the same marginal coverage as before, but better local adaptivity

The interval based on quantile regression is widened if $\hat{q}_{1-\alpha}>0$ and tightened if $\hat{q}_{1-\alpha}<0$

Section 4

Conformal Classification

From trees/forests to probabilities

From the previous lecture, a classification model, e.g., tree, forest, outputs estimated class probabilities

$$\hat{p}_k(x) \approx \mathbb{P}(Y = k \mid X = x), \qquad k = 1, \dots, K$$

- \bullet For a decision tree, these probabilities come from the empirical class proportions in the leaf that contains x
- For a random forest, we average the class proportions over many trees
- For more complex models (logistic regression, neural nets, etc.), the model directly outputs estimated probabilities

Idea for conformal prediction:

We only need the probability vector

$$\hat{f}(x) = (\hat{p}_1(x), \dots, \hat{p}_K(x)),$$

to construct adequate conformal scores and guarantee that the prediction set contains the true class of a new \boldsymbol{x} with high probability

Split conformal classification with likelihood scores

A very common choice of conformity score for classification is

$$s(\hat{f}(x),y) = 1 - \hat{p}_y(x)$$

i.e. small score = high predicted probability for the candidate label y

- ${\bf 0}$ Split data into train D_1 and calibration D_2 and fit \hat{f} on D_1
- On the calibration set, compute conformity scores

$$\underline{S_i} = s\big(\hat{f}(X_i), Y_i\big) = 1 - \hat{p}_{Y_i}(X_i), \quad i \in \mathcal{D}_2$$

(1-likelihood assigned to correct class)

Let

$$\hat{q}_{1-\alpha} = S_{(\lceil (1-\alpha)(n_2+1)\rceil)}$$

be the $\lceil (1-\alpha)(n_2+1) \rceil$ -th smallest score

Split conformal classification with likelihood scores

 \bullet For a new x, define the conformal prediction set

$$\widehat{C}_{\alpha}(x) = \big\{ y \in \mathcal{Y} : s(\widehat{f}(x), y) \leq \widehat{q}_{1-\alpha} \big\}$$

 \Rightarrow same marginal coverage guarantee as in regression with the smallest prediction sets on average

But, it has poor conditional coverage as the same threshold $\hat{q}_{1-\alpha}$ is applied to

- an "easy" point (probability mass on one class)
- a "hard" point (flat probability vector)

Example: moderate classifier

We follow Zaffran's toy example with miscoverage level $\alpha=0.1$ and three labels $\mathcal{Y}=\{\log, \mathrm{tiger}, \mathrm{cat}\}$

For each calibration point (X_i,Y_i) , we have predicted probabilities $(\hat{p}_{\text{dog}}(X_i),\hat{p}_{\text{tiger}}(X_i),\hat{p}_{\text{cat}}(X_i))$, and define the score $S_i=1-\hat{p}_{Y_i}(X_i)$

i	Y_i	\hat{p}_{dog}	\hat{p}_{tiger}	\hat{p}_{cat}	S_i
1	dog	0.95	0.02	0.03	0.05
2	dog	0.90	0.05	0.05	0.10
3	dog	0.85	0.10	0.05	0.15
4	tiger	0.15	0.60	0.25	0.40
5	tiger	0.15	0.55	0.30	0.45
6	tiger	0.20	0.50	0.30	0.50
7	tiger	0.15	0.45	0.40	0.55
8	cat	0.25	0.40	0.35	0.65
9	cat	0.20	0.45	0.35	0.65
10	cat	0.20	0.35	0.45	0.55

 \bullet The split-conformal threshold is $\hat{q}_{1-\alpha} = S_{([(1-\alpha)(n_2+1)])} = S_{(10)} = 0.65$

For a test point with $(p_{\rm dog},p_{\rm tiger},p_{\rm cat})=(0.05,0.60,0.35)$, the scores are

$$S(dog) = 0.95, \quad S(tiger) = 0.40, \quad S(cat) = 0.65$$

Thus,
$$\widehat{C}_{\alpha}(x_{\mathsf{test}}) = \{y : S(y) \leq \widehat{q}_{1-\alpha}\} = \{\mathsf{tiger}, \mathsf{cat}\}$$

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Same example: sharper classifier, smaller sets

i	Y_i	\hat{p}_{dog}	\hat{p}_{tiger}	\hat{p}_{cat}	S_i
1	dog	0.95	0.02	0.03	0.05
2	dog	0.90	0.05	0.05	0.10
3	dog	0.85	0.10	0.05	0.15
4	tiger	0.15	0.60	0.25	0.40
5	tiger	0.15	0.55	0.30	0.45
6	tiger	0.20	0.50	0.30	0.50
7	tiger	0.15	0.45	0.40	0.55
8	cat	0.25	0.40	0.35	0.65
9	cat	0.20	0.45	0.35	0.65
10	cat	0.20	0.35	0.45	0.55

The new threshold is $\hat{q}_{1-\alpha}=S_{([(1-\alpha)(n_2+1)])}=S_{(10)}=0.45$

For the same test point,

$$S(dog) = 0.95, \quad S(tiger) = 0.40, \quad S(cat) = 0.65$$

Now the prediction set is

$$\widehat{C}_{\alpha}\left(x_{\mathsf{test}}\right) = \left\{y : S(y) \leq 0.45\right\} = \left\{\mathsf{tiger}\right\}$$

Better classifier on calibration set yields smaller prediction sets

Adaptive Prediction Sets (Angelopoulos et al., 2021)

Idea: We trade the previous method yielding smallest average size with a new method that adapts to the hardness of the problem and uses predicted probabilities of all classes (not only the true class) to build conformity scores

Let
$$\hat{f}(x) = (\hat{p}_1(x), \dots, \hat{p}_K(x))$$
 be predicted class probabilities

For each point x:

Sort classes by decreasing probability:

$$\hat{p}_{\pi_x(1)}(x) \geq \hat{p}_{\pi_x(2)}(x) \geq \cdots \geq \hat{p}_{\pi_x(K)}(x)$$

 ${\bf 2}$ For each calibration point $(X_i,Y_i),\,i\in D_2,$ compute a conformity score

$$R_i = \sum_{j=1}^{k_i} \hat{p}_{\pi_i(j)}(X_i), \quad \text{ where } \pi_i(k_i) = Y_i$$

This is the cumulative probability of classes at least as likely as the true one

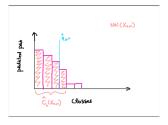
Adaptive Prediction Sets (Angelopoulos et al., 2021)

- **3** Compute $\hat{q}_{1-\alpha}$ the $\lceil (1-\alpha)(n_2+1) \rceil$ -th smallest score
- lacktriangle For a test point x, form

$$\widehat{C}_{\alpha}(x) = \{\pi_x(1), \dots, \pi_x(k_x)\},$$

where k_x is the smallest index such that

$$\sum_{j=1}^{k_x} \hat{p}_{\pi_x(j)}(x) \ge \hat{q}_{1-\alpha}$$



Adaptive Prediction Sets: example

Scores on the calibration set

i	Y_i	\hat{p}_{dog}	\hat{p}_{tiger}	\hat{p}_{cat}	S_i
1	dog	0.95	0.02	0.03	0.95
2	dog	0.90	0.05	0.05	0.90
3	dog	0.85	0.10	0.05	0.85
4	tiger	0.05	0.85	0.10	0.85
5	tiger	0.05	0.80	0.15	0.80
6	tiger	0.05	0.75	0.20	0.75
7	tiger	0.10	0.75	0.15	0.75
8	cat	0.25	0.40	0.35	0.75
9	cat	0.10	0.30	0.60	0.60
10	cat	0.15	0.30	0.55	0.55

- \bullet The split-conformal threshold is $\hat{q}_{1-\alpha}=0.95$
- \bullet For a test point with $(p_{\rm dog}, p_{\rm tiger}, p_{\rm cat}) = (0.05, 0.45, 0.5)$, $k_x = 2$
- \bullet For a test point with $(p_{\rm dog}, p_{\rm tiger}, p_{\rm cat}) = (0.03, 0.95, 0.02)$, $k_x = 1$

Section 5

Beyond Splitting: Full CP and Jackknife+

Full conformal prediction

Idea: The most probable labels Y_{n+1} live in \mathcal{Y} , and have a low enough conformity score. By looping over all possible $y \in \mathcal{Y}$, the ones leading to the smallest conformity scores will be found

For each test point x_{n+1} and candidate label y:

- **1** Augment the dataset with (x_{n+1}, y)
- ② Fit the algorithm on all n+1 points
- Ompute scores for each observation (including the test one)
- lacktriangle Keep y in the prediction set if its score is not too extreme
 - Uses all data both for training and calibration, but
 - \bullet Extremely expensive: re-fit the model for many candidate y values

Full conformal prediction

Finite sample guarantees are obtained under exchangeability (train points and test point) and symmetry of the algorithm

Theorem (Vovk, 2005)

Suppose that

- $(X_i, Y_i)_{i=1}^{n+1}$ are exchangeable, and
- \bullet the algorithm $\mathcal A$ is symmetric (its output depends only on the set of training points, not on their order)

Then, full CP applied on $(X_i,Y_i)_{i=1}^n \cup \{X_{n+1}\}$ outputs $\widehat{C}_{lpha}\left(\cdot\right)$ such that

$$1-\alpha \ \leq \ \mathbb{P}\big(Y_{n+1} \in \hat{C}_{\alpha}(X_{n+1})\big) \ \leq \ 1-\alpha + \frac{1}{n+1},$$

where the upper bound holds if the scores are a.s. distinct

Jackknife and CV variants

- \bullet Compute conformity scores based on LOO predictors $\hat{f}^{-i},$ trained without sample i
- Jackknife+ and CV+ use these LOO (or K-fold) models to produce conformal-like intervals that:
 - re-use data more efficiently than a single split
 - still enjoy marginal coverage guarantees under mild conditions

We will not go into the full formulas here; idea is to connect to the ${\sf CV}$ / bagging ideas you have already seen.

Section 6

Summary

Take-home messages

- Conformal prediction wraps around any black-box predictor
- Under exchangeability, we get finite-sample marginal coverage
- Split CP is simple and robust:
 - use proper training set to get point predictor
 - use calibration set to get residual (or score) quantile
- Choice of score matters:
 - residuals, studentised residuals, CQR scores, likelihood scores for classification
- Classification version connects nicely with probability trees / random forests

References

Introductions and tutorials

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Foundational work

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Methodological developments

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- Barber R., Candès E., Ramdas A. (2021)