Introduction to Conformal Prediction Extension to missing values

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- Standard Split Conformal Prediction for Mean-Regression Improving Adaptiveness: Conformalized Quantile Regression Generalized SCP Framework
- Take-home-messages and open directions
- Quantifying Predictive Uncertainty with Missing Values
- Conclusion

Setting

- $(X, Y) \in \mathbb{R}^d \times \mathbb{R}$ random variables
- *n* training samples $(X^{(i)}, Y^{(i)})_{i=1}^{n}$
- Goal: predict an unseen point Y⁽ⁿ⁺¹⁾ at X⁽ⁿ⁺¹⁾ with confidence
- How? Given a miscoverage level $\alpha \in [0,1]$, build a predictive set \mathcal{C}_{α} such that:

$$\mathbb{P}\left\{Y^{(n+1)} \in \mathcal{C}_{\alpha}\left(X^{(n+1)}\right)\right\} \ge 1 - \alpha, \tag{1}$$

and \mathcal{C}_{lpha} should be as small as possible, in order to be informative

- Construction of the predictive intervals should be
 - agnostic to the model
 - agnostic to the data distribution
 - $\circ~$ valid in finite samples

Standard Split Conformal Prediction for Mean-Regression

Improving Adaptiveness: Conformalized Quantile Regression Generalized SCP Framework

Take-home-messages and open directions

Quantifying Predictive Uncertainty with Missing Values

Split Conformal Prediction (SCP)^{1,2,3}: toy example



¹Vovk et al. (2005), Algorithmic Learning in a Random World ²Papadopoulos et al. (2002), Inductive Confidence Machines for Regression, ECML ³Lei et al. (2018), Distribution-Free Predictive Inference for Regression, JRSS B

Split Conformal Prediction (SCP)^{1,2,3}: training step



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Split Conformal Prediction (SCP)^{1,2,3}: prediction step



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Standard mean-regression SCP: formally

- Split randomly the training data into a proper training set (size #Tr) and a calibration set (size #Cal)
- 2. Train your algorithm on the proper training set to obtain \hat{A}
- 3. On the calibration set, get prediction values with \hat{A}
- 4. Obtain a set of #Cal + 1 conformity scores:

$$\mathcal{S} = \{S^{(i)} = |\hat{A}(X^{(i)}) - Y^{(i)}|, i \in \operatorname{Cal}\} \cup \{+\infty\}$$

(+ worst-case scenario)

- 5. Compute the 1α quantile of these scores, noted $q_{1-\alpha}(\mathcal{S})$
- 6. For a new point $X^{(n+1)}$, output

$$\widehat{C}_{\alpha}\left(X^{(n+1)}\right) = \left[\widehat{A}\left(X^{(n+1)}\right) - q_{1-\alpha}\left(S\right); \widehat{A}\left(X^{(n+1)}\right) + q_{1-\alpha}\left(S\right)\right]$$

SCP theoretical foundation

Definition (Exchangeability)

 $(X^{(i)}, Y^{(i)})_{i=1}^{n}$ are exchangeable if for any permutation σ of $[\![1, n]\!]$ we have:

$$\begin{aligned} & \mathcal{L}\left(\left(X^{(1)}, \, Y^{(1)}\right), \dots, \left(X^{(n)}, \, Y^{(n)}\right)\right) \\ = & \mathcal{L}\left(\left(X^{(\sigma(1))}, \, Y^{(\sigma(1))}\right), \dots, \left(X^{(\sigma(n))}, \, Y^{(\sigma(n))}\right)\right), \end{aligned}$$

where \mathcal{L} designates the joint distribution.

Examples of exchangeable sequences

• i.i.d. samples

 $\bullet\,$ The components of ${\cal N}\,$

$$\begin{pmatrix}m\\\vdots\\\vdots\\m\end{pmatrix}, \begin{pmatrix}\sigma^2&&&\\&\ddots&\gamma^2\\&\gamma^2&\ddots\\&&&\sigma^2\end{pmatrix}\end{pmatrix}$$

SCP enjoys finite sample guarantees proved in Vovk et al. (2005); Lei et al. (2018).

Theorem

Suppose $(X^{(i)}, Y^{(i)})_{i=1}^{n+1}$ are exchangeable (or i.i.d.). SCP applied on $(X^{(i)}, Y^{(i)})_{i=1}^{n}$ outputs $\widehat{C}_{\alpha}(X^{(n+1)})$ such that:

$$\mathbb{P}\left\{Y^{(n+1)}\in\widehat{C}_{lpha}\left(X^{(n+1)}
ight)
ight\}\geq1-lpha.$$

Additionally, if the scores $\{S^{(i)}\}_{i \in Cal}$ are a.s. distinct:

$$\mathbb{P}\left\{Y^{(n+1)} \in \widehat{\mathcal{C}}_{\alpha}\left(X^{(n+1)}\right)\right\} \leq 1 - \alpha + \frac{1}{\#\mathrm{Cal} + 1}.$$

× Marginal coverage: $\mathbb{P}\left\{Y^{(n+1)} \in \widehat{C}_{\alpha}\left(X^{(n+1)}\right) | X^{(n+1)} = x\right\} \ge 1 - \alpha$

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⁴Romano et al. (2019), Conformalized Quantile Regression, NeurIPS



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$$\hookrightarrow \quad S^{(i)} := \max\left\{\widehat{QR}_{\alpha/2}\left(X^{(i)}\right) - Y^{(i)}, Y^{(i)} - \widehat{QR}_{1-\alpha/2}\left(X^{(i)}\right)\right\}$$

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Generalization: SCP is defined by the conformity scores

- Split randomly the training data into a proper training set (size #Tr) and a calibration set (size #Cal)
- 2. Train your algorithm on the proper training set to obtain \hat{A}
- 3. On the calibration set, obtain #Cal + 1 conformity scores

$$\mathcal{S} = \{S^{(i)} = \mathbf{s}\left(X^{(i)}, Y^{(i)}\right), i \in \operatorname{Cal}\} \cup \{+\infty\}$$

Ex 1: $\mathbf{s}(x, y) = |\hat{A}(x) - y|$ in mean-regression with standard scores Ex 2: $\mathbf{s}(x, y) = \max\left(\widehat{QR}_{\alpha/2}(x) - y, y - \widehat{QR}_{1-\alpha/2}(x)\right)$ in CQR

4. Compute the $1 - \alpha$ quantile of these scores, noted $q_{1-\alpha}(S)$ 5. For a new point $X^{(n+1)}$, return

$$\widehat{C}_{\alpha}\left(X^{(n+1)}\right) := \{y \text{ such that } \mathbf{s}\left(\widehat{A}\left(X^{(n+1)}\right), y\right) \leq q_{1-\alpha}\left(\mathcal{S}\right)\}$$

 \hookrightarrow The definition of the conformity scores is crucial, as they incorporate almost all the information: data + underlying model

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ight\}\leq1-lpha+rac{1}{\#\mathrm{Cal}+1}.$$

× Marginal coverage: $\mathbb{P}\left\{Y^{(n+1)} \in \widehat{C}_{\alpha}\left(X^{(n+1)}\right) | X^{(n+1)} = x\right\} \ge 1 - \alpha$

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SCP: summary

Split conformal prediction is simple to compute and works:

- any regression (and classification link to classification) algorithm (neural nets, random forest...);
- distribution-free as long as the data is exchangeable;
- finite sample.

Two interests:

- quantify the uncertainty of the underlying model \hat{A}
- output predictive regions

Note that the theoretical guarantee is **marginal** over the joint distribution of (X, Y), and **not conditional**. That is, there is no guarantee that for any $x \in \mathbb{R}$:

$$\mathbb{P}\left\{Y^{(n+1)} \in \widehat{C}_{\alpha}\left(X^{(n+1)}\right) | X^{(n+1)} = x\right\} \ge 1 - \alpha.$$
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- 1. Providing a form of conditional guarantee
- 2. Tradeoffs between computational cost and statistical efficiency (i.e. variability of the estimators, *efficiency* of the predictive sets)
- 3. Going beyond the exchangeability assumption

CP is a very active field of research. Many developments focus on adapting CP to specific frameworks, such as: Survival Analysis (Candès et al., 2023), Causal Inference (Lei and Candès, 2021; Jin et al., 2023), NLP (Schuster et al., 2022), RL (Taufiq et al., 2022), applications (medical (Angelopoulos et al., 2022; Lu et al., 2022), energy (Kath and Ziel, 2021), etc.) and more.

Quantifying Predictive Uncertainty with Missing Values

Learning with Missing Data

Conformal Prediction with Missing Values

Missing Data Augmentation

Experimental Results

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Experimental Results

Missing values: ubiquitous in data science practice

Y	X_1	X_2	X_3	X_4	X_5	X_6
22.42	0.55	0.67	0.03	0.75	0.05	0.05
8.26	0.72	0.18	0.55	0.05	0.73	0.50
19.41	0.60	0.58	NA	NA	NA	0.40
19.75	0.54	0.43	0.96	0.77	0.06	0.66
-7.32	NA	0.19	NA	0.02	0.83	-0.04-
-13.55	0.65	0.69	0.50	0.15	NA	0.87
20.75	0.43	0.74	0.61	0.72	0.52	0.35
9.26	0.89	NA	0.84	0.01	0.73	NA
<u> 9.68 </u>	0.963	0.45	0.65	0.04	0.06	NA

If each entry has a probability 0.01 of being missing:

 $d=6
ightarrow \, pprox 94\%$ of rows kept

 $d=300
ightarrow\,pprox\,5\%$ of rows kept

One of the **ironies of Big Data** is that missing data play an ever more significant role.⁵

⁵Zhu et al. (2019), High-dimensional PCA with heterogeneous missingness, JRSS B

Handling missing values depends on pattern and mechanism

- $(X, Y) \in \mathbb{R}^d \times \mathbb{R}$ random variables.
- M ∈ {0,1}^d is defined as M_j = 1 ⇔ X_j is missing.
 M is called the mask or the missing pattern.

Example

We observe
$$(NA, 6, 2)(-1, NA, 2)(-1, NA, NA)$$
. Then $m = (1, 0, 0)m = (0, 1, 0)m = (0, 1, 1)$.

There are 2^d patterns (statistical and computational challenges).

Three mechanisms⁶ can generate missing values.
 → Missing Completely At Random (MCAR):
 P(M = m|X) = P(M = m) for all m ∈ {0,1}^d. M ⊥⊥ X,

missingness does not depend on the variables.

⁶Rubin (1976), Inference and missing data, Biometrika

Supervised learning with missing values

Impute-then-regress procedures are widely used.

- 1. Replace NA using an imputation function ϕ (e.g. the mean).
- 2. Train your algorithm (Random Forest, Neural Nets, etc.) on

the imputed data:

$$\left\{\underbrace{\phi\left(X_{obs(M^{(i)})}^{(i)}, M^{(i)}\right)}_{imputed X^{(i)}}, Y^{(i)}\right\}_{k=1}^{n}$$

 \checkmark : Le Morvan et al. (2021)⁷ show that for any deterministic imputation and universal learner this procedure is Bayes-consistent.

✗: Ayme et al. (2022)⁸ show that even for very simple distributions (linear model, Gaussian noise), may suffer from curse of dimensionality.

⁷ Le Morvan et al. (2021), What's a good imputation to predict with missing values?, NeurIPS ⁸ Ayme et al. (2022), Near-optimal rate of consistency for linear models with missing values, ICML

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Impute-then-regress+conformalization is marginally valid

To apply conformal prediction we need **exchangeable** data.

Lemma (Exchangeability after imp., Zaffran et al., 2023) Assume $(X^{(i)}, M^{(i)}, Y^{(i)})_{i=1}^{n}$ are *i.i.d.* (or exchangeable).

Then, for any missing mechanism, for almost all imputation function ϕ :

 $\left(\phi\left(X_{obs(M^{(i)})}^{(i)}, M^{(i)}\right), Y^{(i)}\right)_{i=1}^{n}$ are exchangeable.

 \Rightarrow Conformal prediction applied on an imputed data set still enjoys marginal guarantees $^9:$

$$\mathbb{P}\left(Y \in \widehat{C}_{\alpha}\left(X_{\mathsf{obs}(M)}, M\right)\right) \geq 1 - \alpha.$$

Even if the imputation is not accurate, the guarantee will hold.

⁹The upper bound also holds under continuously distributed scores.

CQR performances on an illustrative example

$$Y = \beta^T X + \varepsilon,$$

with $\beta = (1, 2, -1)^T$, $\varepsilon \perp X$ and X and ε are Gaussian.



suffer from high disparities depending on the missing patterns.

Theoretical study of the Gaussian linear model ($Y = \beta^T X + \varepsilon$) generalizes:

Proposition (Oracle intervals under the Gaussian lin. mod.)

$$\mathcal{L}^*_{\alpha}(\textit{m}) = 2 \times q_{1-\alpha/2}^{\mathcal{N}(0,1)} \times \sqrt{\beta_{\mathrm{mis}(\textit{m})}^{\mathcal{T}} \Sigma_{\mathrm{mis}|\mathrm{obs}}^{\textit{m}} \beta_{\mathrm{mis}(\textit{m})} + \sigma_{\varepsilon}^2}$$

- Even with an homoskedastic noise, missingness generates heteroskedasticity
- The uncertainty increases when missing values are associated with larger regression coefficients (i.e. the most predictive variables)

Goal: for any $m \in \mathcal{M} \subset \{0,1\}^d$:

$$\mathbb{P}\left(Y \in \widehat{\mathcal{C}}_{\alpha}\left(X_{\mathsf{obs}(M)}, M\right) | M = m\right) \geq 1 - \alpha.$$

Motivation: equity, first-step-towards-conditional.

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¹⁰Romano et al. (2020), With Malice Toward None: Assessing Uncertainty via Equalized Coverage, Harvard Data Science Review

Missing data augmentation of the calibration set



$$\hookrightarrow S^{(i)} := \max\left\{\widehat{QR}_{\alpha/2}\left(\tilde{X}^{(i)}\right) - Y^{(i)}, Y^{(i)} - \widehat{QR}_{1-\alpha/2}\left(\tilde{X}^{(i)}\right)\right\}$$

CQR-MDA with exact masking in words

- 1. Split the training set into a proper training set and calibration set
- 2. Train the imputation function on the proper training set
- 3. Impute the proper training set
- 4. Train the quantile regressors on the imputed proper training set
- 5. For a test point $(X^{(n+1)}, M^{(n+1)})$:
 - 5.1 For each $j \in \llbracket 1, d \rrbracket$ s.t. $M_j^{(n+1)} = 1$, set $\tilde{M}_i^{(i)} = 1$ for i in Cal s.t. $M^{(i)} \subset M^{(n+1)}$
 - 5.2 Impute the new calibration set
 - 5.3 Compute the calibration correction, i.e. $q_{1-\alpha}(S)$
 - 5.4 Impute the test point
 - 5.5 Predict with the quantile regressors and the correction previously obtained, $q_{1-\alpha}(S)$





Theorem (Zaffran et al., 2023)

If the data is exchangeable and MCAR, then for almost all imputation function the proposed methodology is such that for any $m \in \{0,1\}^d$:

$$\mathbb{P}\left(Y \in \widehat{\mathcal{C}}_{\alpha}\left(X_{obs(M)}, M\right) | M = m\right) \geq 1 - \alpha,$$

and if additionally the scores are almost surely distinct:

$$\mathbb{P}\left(Y \in \widehat{\mathcal{C}}_{\alpha}\left(X_{obs(M)}, M\right) | M = m\right) \leq 1 - \alpha + \frac{1}{1 + \# \mathrm{Cal}^{\mathrm{m}}}.$$

Empirical coverages



Empirical lengths



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- Imputation by iterative ridge (\sim conditional expectation)
- Concatenate the mask in the features
- Neural network, fitted to minimize the pinball loss
- (Semi)-synthetic experiments:
 - $\circ~$ MCAR missing values, with probability 20%
 - \circ 100 repetitions

Synthetic experiments (Gaussian linear model, d = 10)





- $igstarrow: ext{marginal coverage, i.e.} \ \mathbb{P}(Y \in \hat{C}_lpha(X,M))$
- $oldsymbol{
 abla}$: lowest coverage, i.e. $\min_{m\in\mathcal{M}}\mathbb{P}(Y\in\hat{C}_{lpha}(X,m)|M=m)$
- $\begin{array}{l} \blacktriangle : \text{highest coverage, i.e.} \\ \max_{m \in \mathcal{M}} \mathbb{P}(Y \in \hat{C}_{\alpha}(X,m) | M = m) \end{array}$

Semi-synthetic experiments



- 30 hospitals
- More than 30 000 trauma patients
- 4 000 new patients per year
- 250 continuous and categorical variables
 - $\hookrightarrow \mathsf{Many} \text{ useful statistical tasks}$

Predict the level of platelets upon arrival at hospital, given 7 covariates chosen by medical doctors.

These covariates are not always observed: from 0% to 24% of missing values by features, with a total average of 7%.



Introduction to (Split) Conformal Prediction

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Conclusion

- Consistency of universal quantile learner when chained with almost any imputation function.
- CP-MDA-Nested (link to CP-MDA-Nested), an algorithm which does not discard any calibration point.



- CP marginal guarantees hold on the imputed data set.
- Missingness introduces additional heteroskedasticity, creating a need for quantile regression based methods.
- CQR fails to attain coverage conditional on the missing pattern.
- Missing data augmentation is the first method to output predictive intervals with missing values.
- Missing data augmentation attains conditional coverage with respect to the missing pattern (in MCAR setting).

Thank you!

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Appendix

Quantile regression



Warning

No theoretical guarantee with a finite sample

$$\mathbb{P}\left(Y\in\left[\hat{Q}_{Y|X}(eta/2);\hat{Q}_{Y|X}(1-eta/2)
ight]
ight)
eq1-eta$$

SCP: what choices for the regression scores?

	Standard SCP	Locally weighted SCP	CQR				
	Vovk et al. (2005)	Lei et al. (2018)	Romano et al. (2019)				
s(X Y)	$ \hat{A}(X) - Y $	$ \hat{A}(X)-Y $	$\max(\widehat{QR}_{\alpha/2}(X) - Y,$				
5 (7(, 1)		$\hat{\rho}(X)$	$Y - \widehat{QR}_{1-\alpha/2}(X))$				
$\hat{C}(\mathbf{v})$	$\left[\hat{A}(w) \pm \sigma_{1}(w)\right]$	$\left[\hat{A}(y) \pm \sigma - (S)\hat{S}(y)\right]$	$[\widehat{QR}_{\alpha/2}(x)-q_{1-\alpha}(\mathcal{S});$				
$C_{\alpha}(x)$	$\left[A(x) \pm q_{1-\alpha}(0)\right]$	$\left[A(x) \pm q_{1-\alpha}(O)p(x)\right]$	$\widehat{QR}_{1-\alpha/2}(x) + q_{1-\alpha}(\mathcal{S})]$				
Visu.	0 2 4 X	0 2 4 X	0 2 4 X				
\checkmark	black-box around a	black-box around a	adaptive				
	"usable" prediction	"usable" prediction					
×	not adaptive	limited adaptiveness	no black-box around a				
			"usable" prediction				

SCP in classification (from C. Boyer and M. Zaffran tutorial)

- $Y^{(i)} \in \{1, \dots, C\}$ (C classes)
- $\hat{A}(X) = (\hat{p}_1(X), \dots, \hat{p}_C(X))$ (estimated probabilities)
- Score of the *i*-th calibration point: $S^{(i)} = 1 (\hat{A}(X^{(i)}))_{Y^{(i)}}$
- For a new point $X^{(n+1)}$, return

$$\widehat{C}_{\alpha}\left(X^{(n+1)}\right) = \{y \text{ such that } s(\widehat{A}\left(X^{(n+1)}\right), y) \leq q_{1-\alpha}\left(\mathcal{S}\right)\}$$

SCP in classification in practice

Ex: $Y^{(i)} \in \{$ "dog", "tiger", "cat" }, with $\alpha = 0.1$

• Scores on the calibration set

$\operatorname{Cal}^{(i)}$	-(2)	1.2		- Charles	and the second second		S.V.	1	R	200
$\hat{p}_{dog}\left(X^{(i)} ight)$	0.95	0.90	0.85	0.15	0.15	0.20	0.15	0.15	0.25	0.20
$\hat{p}_{tiger}\left(X^{(i)}\right)$	0.02	0.05	0.10	0.60	0.55	0.50	0.45	0.40	0.35	0.45
$\hat{p}_{cat}\left(X^{(i)}\right)$	0.03	0.05	0.05	0.25	0.30	0.30	0.40	0.45	0.40	0.35
S (<i>i</i>)	0.05	0.1	0.15	0.40	0.45	0.50	0.55	0.55	0.6	0.65

•
$$q_{1-\alpha}(S) = 0.65$$

• $\hat{A}(X^{(n+1)}) = (0.05, 0.60, 0.35)$
 $\hookrightarrow s(\hat{A}(X^{(n+1)}), \text{"dog"}) = 0.95$
 $\hookrightarrow s(\hat{A}(X^{(n+1)}), \text{"tiger"}) = 0.40 \le q_{1-\alpha}(S)$
"tiger" $\in \hat{C}_{\alpha}(X^{(n+1)})$
 $\hookrightarrow s(\hat{A}(X^{(n+1)}), \text{"cat"}) = 0.65 \le q_{1-\alpha}(S)$ "cat" $\in \hat{C}_{\alpha}(X^{(n+1)})$
• $\hat{C}_{\alpha}(X^{(n+1)}) = \{\text{"tiger", "cat"}\}$

SCP in classification in practice

Ex: $Y^{(i)} \in \{$ "dog", "tiger", "cat" }, with $\alpha = 0.1$

• Scores on the calibration set

$\operatorname{Cal}^{(i)}$		1.2		\$	- Allen-			1		sk.
$\hat{p}_{dog}(X^{(i)})$	0.95	0.90	0.85	0.05	0.05	0.05	0.05	0.10	0.10	0.15
$\hat{p}_{tiger}(X^{(i)})$	0.02	0.05	0.10	0.85	0.80	0.75	0.70	0.25	0.30	0.30
$\hat{p}_{cat}\left(X^{(i)}\right)$	0.03	0.05	0.05	0.10	0.15	0.20	0.25	0.65	0.60	0.55
S (<i>i</i>)	0.05	0.1	0.15	0.15	0.20	0.25	0.30	0.35	0.40	0.45

•
$$q_{1-\alpha}(S) = 0.45$$

• $\hat{A}(X^{(n+1)}) = (0.05, 0.60, 0.35)$
 $\hookrightarrow s(\hat{A}(X^{(n+1)}), \text{"dog"}) = 0.95$
 $\hookrightarrow s(\hat{A}(X^{(n+1)}), \text{"tiger"}) = 0.40 \le q_{1-\alpha}(S)$
"tiger" $\in \hat{C}_{\alpha}(X^{(n+1)})$
 $\hookrightarrow s(\hat{A}(X^{(n+1)}), \text{"cat"}) = 0.65$
"cat" $\notin \hat{C}_{\alpha}(X^{(n+1)})$
• $\hat{C}(X^{(n+1)}) = \xi$ "tiger" ξ

- Facts about the previous method
 - $\circ\;$ prediction sets with the smallest average size
 - $\circ~$ undercover hard subgroups
 - $\circ~$ overcover easy ones
- Other types of scores can be used to improve the conditional coverage (as in regression with CQR or localized)

SCP in classification: Adaptive Prediction Sets

- 1. Sort in decreasing order $\hat{p}_{\sigma_i(1)}\left(X^{(i)}\right) \geq \ldots \geq \hat{p}_{\sigma_i(C)}\left(X^{(i)}\right)$
- 2. $S^{(i)} = \sum_{k=1}^{\sigma_i^{-1}(Y^{(i)})} \hat{p}_{\sigma_i(k)}(X^{(i)}) \qquad (\text{sum of the estimated probabilities associated})$

to classes at least as large as that of the true class Y_i)

3. Return the classes $\sigma^{(n+1)}(1),\ldots,\sigma^{(n+1)}(r^{\star})$ where

$$r^{\star} = \operatorname*{arg\,max}_{1 \leq r \leq C} \left\{ \sum_{k=1}^{r} \hat{p}_{\sigma^{(n+1)}(k)} \left(X^{(n+1)} \right) < q_{1-\alpha}(\mathcal{S}) \right\} + 1$$



Ex: $Y_i \in \{$ "dog", "tiger", "cat" $\}$, with $\alpha = 0.1$

• Scores on the calibration set

$\operatorname{Cal}^{(i)}$		1.	0	s)-				5		94¢).
$\hat{p}_{dog}\left(X^{(i)}\right)$	0.95	0.90	0.85	0.05	0.05	0.05	0.10	0.25	0.10	0.15
$\hat{p}_{tiger}\left(X^{(i)} ight)$	0.02	0.05	0.10	0.85	0.80	0.75	0.75	0.40	0.30	0.30
$\hat{p}_{cat}\left(X^{(i)} ight)$	0.03	0.05	0.05	0.10	0.15	0.20	0.15	0.35	0.60	0.55
S ⁽ⁱ⁾	0.95	0.90	0.85	0.85	0.80	0.75	0.75	0.75	0.60	0.55

- $q_{1-\alpha}(\mathcal{S}) = 0.95$
- Ex 1: $\hat{A}(X^{(n+1)}) = (0.05, 0.45, 0.5), r^* = 2$ $\hat{C}_{\alpha}(X^{(n+1)}) = \{\text{"tiger", "cat"}\}$ • Ex 2: $\hat{A}(X^{(n+1)}) = (0.03, 0.95, 0.02), r^* = 1$ $\hat{C}_{\alpha}(X^{(n+1)}) = \{\text{"tiger"}\}$
Jackknife/cross-val (from C. Boyer and M. Zaffran tutorial)

Beyond the limitations of SCP

- SCP is computationally attractive: it only requires fitting the model one time
- Problem: it sacrifices statistical efficiency
 - requiring splitting the data into training and calibration datasets
- → Full (or transductive) conformal prediction
 - $\circ~$ avoids data splitting
 - $\circ\;$ at the cost of many more model fits
 - Historically, full conformal prediction was developed first
 - Idea: we know that the true label Y⁽ⁿ⁺¹⁾ lives somewhere in 𝔅 so if we loop over all possible y ∈ 𝔅, then we will eventually hit the data point (X⁽ⁿ⁺¹⁾, Y⁽ⁿ⁺¹⁾), which is statistically plausible with the first n data points
 - Hence the name as full conformal prediction directly computes this loop

Method: for a candidate $(X^{(n+1)}, y)$,

1. Get \hat{A}_{y} by training on $\{(X^{(1)}, Y^{(1)}), \dots, (X^{(n)}, Y^{(n)})\} \cup \{(X^{(n+1)}, y)\}$

2. Scores

$$\mathcal{S} = \left\{ s(\hat{A}_{y}\left(X^{(i)}, Y^{(i)}\right) \right\} \cup \left\{ s(\hat{A}_{y}\left(X^{(n+1)}\right), y) \right\}$$

3. $y \in \widehat{C}_{\alpha}\left(X^{(n+1)}\right)$ if $s(\widehat{A}_{y}\left(X^{(n+1)}\right), y) \leq q_{1-\alpha}(\mathcal{S})$

- Theoretical guarantees (provided that the learning algorithm handles exchangeable training data in a symmetric way)
- X Computationally costly: not used in practice

Other methods for conformal prediction



Jackknife: naive predictive interval

• Based on leave-one-out (LOO) residuals



- $\mathcal{D}^n = \left\{ \left(X^{(1)}, Y^{(1)} \right), \dots, \left(X^{(n)}, Y^{(n)} \right) \right\}$ training data
- Get \hat{A}^{-i} by training on $\mathcal{D}^n \setminus \left(X^{(i)}, Y^{(i)}
 ight)$
- LOO scores $S = \left\{ \left| \hat{A}^{-i} \left(X^{(i)} \right) Y^{(i)} \right| \right\}_i \cup \{+\infty\}$ (in standard reg)
- Get \hat{A} by training on \mathcal{D}^n
- Build the predictive interval: $\left[\hat{A}\left(X^{(n+1)}
 ight)\pm q_{1-lpha}(\mathcal{S})
 ight]$

Warning

No guarantee on the prediction of \hat{A} with scores based on $(\hat{A}^{-i})_i$

Jackknife+ (Barber et al., 2021b)

Based on leave-one-out (LOO) residuals



- $\mathcal{D}^n = \left\{ \left(X^{(1)}, Y^{(1)} \right), \dots, \left(X^{(n)}, Y^{(n)} \right) \right\}$ training data
- Get \hat{A}^{-i} by training on $\mathcal{D}^n \setminus \left(X^{(i)}, Y^{(i)}
 ight)$
- LOO predictions (in standard reg) $S_{up/down} = \left\{ \hat{A}^{-i} \left(X^{(n+1)} \right) \pm | \hat{A}^{-i} \left(X^{(i)} \right) - Y^{(i)} | \right\}_{i} \cup \{\pm \infty\}$
- Build the predictive interval: $\left[q_{\alpha/2}(\mathcal{S}_{\mathsf{down}}); q_{1-\alpha/2}(\mathcal{S}_{\mathsf{up}})\right]$

Theorem

If $\mathcal{D}^n \cup (X^{(n+1)}, Y^{(n+1)})$ are exchangeable and the algorithm treats the data points symmetrically, then $\mathbb{P}(Y^{(n+1)} \in \widehat{C}_{\alpha}(X^{(n+1)})) \geq 1 - 2\alpha$.

CV+ (Barber et al., 2021b)



- Based on cross-validation residuals
- $\mathcal{D}^n = \left\{ \left(X^{(1)}, Y^{(1)} \right), \dots, \left(X^{(n)}, Y^{(n)} \right) \right\}$ training data
- 1. Split \mathcal{D}^n into K folds F_1, \ldots, F_K
- 2. Get \hat{A}^{-F_k} by training on $\mathcal{D}^n \setminus F_k$
- 3. Cross-val predictions (in standard reg) $\mathcal{S}_{up/down} = \left\{ \left\{ \hat{A}^{-F_k} \left(X^{(n+1)} \right) \pm | \hat{A}_{-F_k} \left(X^{(i)} \right) - Y^{(i)} | \right\}_{i \in F_k} \right\}_k \cup \{ \pm \infty \}$
- 4. Build the predictive interval: $[q_{\alpha}(\mathcal{S}_{down}); q_{1-\alpha}(\mathcal{S}_{up})]$

Theorem

Under data exchangeability and algorithm symmetry, then $\mathbb{P}(Y^{(n+1)} \in \widehat{C}_{\alpha}(X^{(n+1)})) \geq 1 - 2\alpha - \min\left(\frac{2(1-1/K)}{n/K+1}, \frac{1-K/n}{K+1}\right) \geq 1 - 2\alpha - \sqrt{2/n}.$

CP-MDA-Nested

CP-MDA-Exact reminder







CQR-MDA with nested masking in words

1. For a test point $(X^{(n+1)}, M^{(n+1)})$:

1.1 Set
$$\tilde{M}^{(i)} = \max(M^{(i)}, M^{(n+1)})$$
 for *i*
in the calibration set

3 NA NA 1

$ ilde{x}^{(1)}$	-1	NA	NA	1
$ ilde{x}^{(2)}$	4	NA	NA	2
$ ilde{x}^{(3)}$	5	NA	NA	NA
$ ilde{x}^{(4)}$	0	NA	NA	1

- 1.2 Impute the new calibration set
- 1.3 For each augmented calibration point *i*:

1.3.1 Get its score $S^{(i)}$

Impute-then-predict on the augmented

1.3.2 test point $(X^{(n+1)}, \tilde{M}^{(i)})$, giving: $\widehat{QR}_{\alpha/2}(\tilde{X}^{(n+1),i})$ and $\widehat{QR}_{1-\alpha/2}(\tilde{X}^{(n+1),i})$

3	NA	NA	1
3	NA	NA	1
3	NA	NA	NA
3	NA	NA	1

1.3.3 Compute the corrected prediction interval: $[\widehat{QR}_{\alpha/2}(\tilde{X}^{(n+1),i}) - S^{(i)}; \widehat{QR}_{1-\alpha/2}(\tilde{X}^{(n+1),i}) + S^{(i)}] := [Z_{inf}^{(i)}, Z_{sup}^{(i)}]$

1.4 Compute the quantiles $q_{\alpha}(\{Z_{inf}^{(i)}\}_{i \in Cal})$ and $q_{1-\alpha}(\{Z_{sup}^{(i)}\}_{i \in Cal})$ 1.5 Predict $[q_{\alpha}(\{Z_{inf}^{(i)}\}_{i \in Cal}); q_{1-\alpha}(\{Z_{sup}^{(i)}\}_{i \in Cal})]$

Summary of CP-MDA



Towards asymptotic individualized coverage

Consistency of a universal quantile learner after imputation

Let Φ be an imputation function chosen by the user.

Denote

$$g^*_{\beta,\Phi} \in \operatorname*{argmin}_{\substack{g: \mathbb{R}^d \to \mathbb{R}}} \mathbb{E}\left[
ho_{eta}(Y - \underline{g} \circ \Phi(X_{\mathrm{obs}(\mathrm{M})}, M)) \right] := \mathcal{R}_{eta,\phi}(g).$$

Comparison with: argmin $\mathbb{E}\left[\rho_{\beta}(Y - f(X_{obs(M)}, M))\right]$ (informal).

Proposition (Pinball-consistency of an universal learner)

For almost all \mathcal{C}^{∞} imputation function Φ , the function $g^*_{\beta,\Phi} \circ \Phi$ is Bayes optimal for the pinball-risk of level β .

 \hookrightarrow any universally consistent algorithm for quantile regression trained on the data imputed by Φ is pinball-Bayes-consistent.

This is an extension of the result of Le Morvan et al. (2021).

Corollary

For any missing mechanism, for almost all C^{∞} imputation function Φ , if $F_{Y|(X_{obs(M)},M)}$ is continuous, a universally consistent quantile regressor trained on the imputed data set yields asymptotic conditional coverage.

 $\hookrightarrow \mathbb{P}(Y \in \widehat{C}_{\alpha}(x) | X = x, M = m) \ge 1 - \alpha$ for any $m \in \mathcal{M}$ and any $x \in \mathbb{R}^d$, asymptotically with a super quantile learner.

$$(X, Y) \in \mathbb{R}^3 \times \mathbb{R}.$$

$$Y = \beta X + \varepsilon$$

with $\varepsilon \sim \mathcal{N}(0, 1), \ \beta = (1, 2, -1) \text{ and}$

$$(X_1, X_2, X_3) \sim \mathcal{N}\left(\begin{pmatrix} 1\\1\\1\\1 \end{pmatrix}, \begin{pmatrix} 1 & 0.8 & 0.8\\0.8 & 1 & 0.8\\0.8 & 0.8 & 1 \end{pmatrix}\right).$$

All components of X each have a probability 0.2 of being missing, Completely At Random.

- Method: CQR
- Basemodel: neural network
- 200 repetitions
 - \circ train size of 250 points
 - $\circ\,$ calibration size of 250 points
 - \circ test size of 2000 points

d = 10, with missing data augmentation

$$(X, Y) \in \mathbb{R}^{10} \times \mathbb{R}.$$

$$Y = \beta X + \varepsilon$$

with $\varepsilon \sim \mathcal{N}(0, 1), \ \beta = (1, 2, -1, 3, -0.5, -1, 0.3, 1.7, 0.4, -0.3)$
and $(X_1, \dots, X_{10}) \sim \mathcal{N}\left(\begin{pmatrix} 1\\ \vdots\\ \vdots\\ 1 \end{pmatrix}, \begin{pmatrix} 1 & 0.8 & \cdots & 0.8\\ 0.8 & \ddots & \ddots & \vdots\\ \vdots & \ddots & \ddots & 0.8\\ 0.8 & \cdots & 0.8 & 1 \end{pmatrix}\right).$

All components of X each have a probability 0.2 of being missing, Completely At Random.

- Method: CQR
- Basemodel: neural network
- Imputation: iterative (pprox conditional expectation)
- Mask as features: yes
- 100 repetitions
 - train size of 500 points
 - $\circ\,$ calibration size of 250 points
 - \circ test size of 100 points for each pattern size, and 2000 for the marginal test set

Results per pattern size



- Method: CQR
- Basemodel: neural network
- Imputation: iterative (\approx conditional expectation)
- Mask as features: yes
- 100 repetitions
 - \circ train size varies
 - $\circ\,$ calibration size of 1000 points
 - \circ test size of 2000 points

Results on the worst group



Results on the best group





MNAR self masked missingness



MNAR quantile censorship missingness













- 30 hospitals
- More than 30 000 trauma patients
- 4 000 new patients per year
- 250 continuous and categorical variables
 - $\hookrightarrow \mathsf{Many} \text{ useful statistical tasks}$

Predict the level of platelets upon arrival at hospital, given 7 covariates chosen by medical doctors.

These covariates are not always observed.
Data set description i

- Age: the age of the patient (no missing values);
- Lactate: the conjugate base of lactic acid, upon arrival at the hospital (17.66% missing values);
- Delta_hemo: the difference between the hemoglobin upon arrival at hospital and the one in the ambulance (23.82% missing values);
- VE: binary variable indicating if a Volume Expander was applied in the ambulance. A volume expander is a type of intravenous therapy that has the function of providing volume for the circulatory system (2.46% missing values);
- RBC: a binary index which indicates whether the transfusion of Red Blood Cells Concentrates is performed (0.37% missing values);

- SI: the shock index. It indicates the level of occult shock based on heart rate (HR) and systolic blood pressure (SBP), that is SI = $\frac{HR}{SBP}$, upon arrival at hospital (2.09% missing values);
- HR: the heart rate measured upon arrival of hospital (1.62% missing values).

Results with CQR-MDA-Nested

