An introduction to conformal prediction and distribution-free inference (Lecture 3)

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Topics — Lecture 3

Cross-validation based methods

- Challenges for distribution-free CV
- Jackknife & jackknife+
- Theory for jackknife+
- CV+ and cross-conformal
- Theory for cross-conformal (and CV+)
- Connection to algorithmic stability

Summary

Summary & preview

Cross-validation based methods

Challenges for distribution-free CV

Returning to cross-validation

Summarizing different methods:

- Split CP fits $\widehat{\mu}$ to part of the data \leadsto distrib.-free theory
- Full CP: use all the data for $\widehat{\mu}$ and achieves distrib.-free theory, but computationally very expensive
- Can cross-validation based methods offer a compromise?

Returning to cross-validation

Why does distribution-free theory hold for split CP but not for CV?

$$\mathcal{C}(X_{n+1}) = \widehat{\mu}(X_{n+1}) \pm \widehat{q} \implies \text{coverage if } \underbrace{|Y_{n+1} - \widehat{\mu}(X_{n+1})|}_{=R_{n+1}} \le \widehat{q}$$

ullet For split conformal, \widehat{q} is quantile of calibration residuals

$$R_i = |Y_i - \widehat{\mu}(X_i)|, i = n_0 + 1, \dots, n_i$$

and $\widehat{\mu}$ is pretrained $\Rightarrow R_{n_0+1}, \dots, R_n, R_{n+1}$ are exchangeable

ullet For CV, \widehat{q} is quantile of leave-one-out residuals

$$R_i = |Y_i - \widehat{\mu}_{-i}(X_i)|, i = 1, ..., n$$

 $\Rightarrow R_1, \dots, R_n, R_{n+1}$ are not exchangeable

Cross-validation based methods

Cross-validation based methods

Jackknife & jackknife+

Recall leave-one-out CV (also known as jackknife):

$$C(X_{n+1}) = \widehat{\mu}(X_{n+1}) \pm \mathsf{Quantile}_{1-\alpha}(R_i)$$

where $R_i = |Y_i - \widehat{\mu}_{-i}(X_i)|$ are the leave-one-out residuals

- ullet In practice, generally we see pprox 1-lpha coverage
- In theory, coverage may be zero! e.g., $\mathcal A$ runs least squares for n even, or neural net for n odd

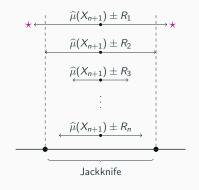
Jackknife can equivalently be defined as:

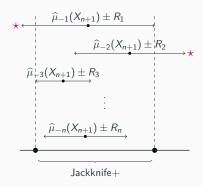
$$\begin{split} \mathcal{C}(X_{n+1}) &= \left[\text{Quantile}_{\mathcal{A}}(\hat{\mu}(X_{n+1}) / / / R_i), \text{Quantile}_{1-\alpha}(\hat{\mu}(X_{n+1}) + R_i) \right] \\ &- \text{Quantile}_{1-\alpha}(-\hat{\mu}(X_{n+1}) + R_i) \end{split}$$

A modified version of the method: the jackknife $+^1$.

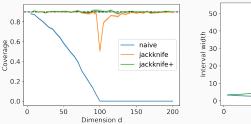
$$\begin{split} \mathcal{C}(X_{n+1}) &= \Big[- \mathsf{Quantile}_{(1-\alpha)(1+1/n)} \big(- \widehat{\mu}_{-i}(X_{n+1}) + R_i \big), \\ &\qquad \qquad \mathsf{Quantile}_{(1-\alpha)(1+1/n)} \big(\widehat{\mu}_{-i}(X_{n+1}) + R_i \big) \Big] \end{split}$$

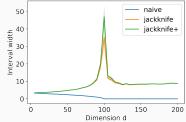
¹B., Candès, Ramdas, Tibshirani 2019, Predictive inference with the jackknife+





Empirical comparison (linear regression with n = 100):





- "Ridgeless" regression minimum- ℓ_2 -norm solution, if d>n
- Note: ridgeless regression is stable except the $d \approx n$ regime²

²Hastie et al 2022, Surprises in High-Dimensional Ridgeless Least Squares Interpolation

Cross-validation based methods

Cross-validation based methods

Theory for jackknife+

Jackknife+ coverage guarantee

Theorem: coverage for jackknife+

If Z_1, \ldots, Z_{n+1} are exchangeable, and $\mathcal A$ is symmetric, then jackknife+ satisfies

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}(X_{n+1})\right\} \ge 1 - 2\alpha$$

(In contrast, jackknife may have zero coverage, in the worst case)

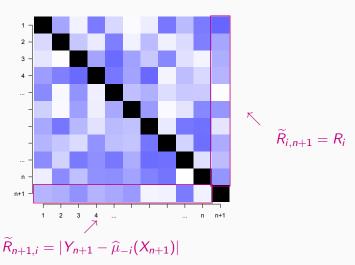
Step 1: a matrix of residuals

For each $i \neq j \in \{1, \dots, n+1\}$ define

$$\widetilde{\mu}_{-ij} = \mathcal{A}(Z_1, \dots, Z_{i-1}, Z_{i+1}, \dots, Z_{j-1}, Z_{j+1}, \dots, Z_{n+1})$$

and define residuals, $\widetilde{R}_{ij} = |Y_i - \widetilde{\mu}_{-ij}(X_i)|$

Matrix of residuals $\widetilde{R} \in \mathbb{R}^{(n+1) \times (n+1)}$



Exchangeability of $\widetilde{R} \in \mathbb{R}^{(n+1)\times(n+1)}$:

- The n(n+1) nondiagonal entries of \widetilde{R} are *not* exchangeable: e.g., expect higher correlation for $\widetilde{R}_{1,2}\&\widetilde{R}_{1,3}$ than $\widetilde{R}_{1,2}\&\widetilde{R}_{3,4}$
- However, \widetilde{R} satisfies $\widetilde{R} \stackrel{\mathrm{d}}{=} \Pi \widetilde{R} \Pi^{\top}$ for any permutation matrix Π

Proof that $\widetilde{R} \stackrel{d}{=} \Pi \widetilde{R} \Pi^{\top}$

Let σ be the permutation corresponding to Π .

Let $f: (\mathcal{X} \times \mathcal{Y})^{n+1} \to \mathbb{R}^{(n+1) \times (n+1)}$ map data to \widetilde{R} :

$$(f(Z_1,\ldots,Z_{n+1}))_{ij}=|Y_i-[\mathcal{A}((Z_k)_{k\neq i,j})](X_i)|$$

Then

$$\widetilde{R} = f(Z_1, \dots, Z_{n+1}) \stackrel{\mathrm{d}}{=} f(Z_{\sigma(1)}, \dots, Z_{\sigma(n+1)}) = \prod \widetilde{R} \Pi^{\top}.$$
exchangeability of the data symmetry of \mathcal{A}

Step 2: relate \widetilde{R} to the jackknife+

$$\begin{aligned} Y_{n+1} &\not\in \mathcal{C}(X_{n+1}), \text{ i.e., jackknife} + \text{ fails to cover} \Longleftrightarrow \\ &\text{either } \underbrace{Y_{n+1} > \text{Quantile}_{(1-\alpha)(1+1/n)} \big(\widehat{\mu}_{-i}(X_{n+1}) + R_i \big)}_{\text{or } Y_{n+1} < -\text{Quantile}_{(1-\alpha)(1+1/n)} \big(-\widehat{\mu}_{-i}(X_{n+1}) + R_i \big)} \\ &\Leftrightarrow \sum_{i=1}^n \underbrace{\mathbb{I}\left\{Y_{n+1} > \widehat{\mu}_{-i}(X_{n+1}) + R_i\right\}}_{\text{same}} \geq n \cdot (1-\alpha)(1+1/n) \end{aligned} \\ &= \mathbb{I}\left\{Y_{n+1} - \widehat{\mu}_{-i}(X_{n+1}) > \widetilde{R}_{i,n+1}\right\} \\ &\leq \mathbb{I}\left\{\widetilde{R}_{n+1,i} > \widetilde{R}_{i,n+1}\right\} \end{aligned}$$

Summary of step 2:

$$Y_{n+1} \notin \mathcal{C}(X_{n+1}) \implies \sum_{i=1}^n \mathbb{1}\left\{\widetilde{R}_{n+1,i} > \widetilde{R}_{i,n+1}\right\} \ge (1-\alpha)(n+1)$$

Combining step 1 & step 2:

we need to show that exchangeability of \widetilde{R} implies

$$\mathbb{P}\left\{\sum_{i=1}^{n}\mathbb{1}\left\{\widetilde{R}_{n+1,i}>\widetilde{R}_{i,n+1}\right\}\geq (1-\alpha)(n+1)\right\}\leq 2\alpha$$

Step 3: the tournament matrix

Consider a tournament played between n + 1 teams:

$$A_{ij} = \mathbb{1}\left\{\widetilde{R}_{ij} > \widetilde{R}_{ji}\right\} \text{ for } i \neq j$$

(and $A_{ii} = 0$)

• $A \stackrel{d}{=} \Pi A \Pi^{\top}$ for perm. matrix Π (inherits this property from \widetilde{R})

•
$$Y_{n+1} \notin \mathcal{C}(X_{n+1}) \Longrightarrow \underbrace{\sum_{i} A_{n+1,i}}_{\text{# games won by team } n+1} \ge (1-\alpha)(n+1)$$

$$\mathbb{P}\left\{Y_{n+1} \not\in \mathcal{C}(X_{n+1})\right\} \leq \mathbb{P}\left\{\sum_{i=1}^{n+1} A_{n+1,i} \geq (1-\alpha)(n+1)\right\}$$

$$= \frac{1}{n+1} \sum_{j=1}^{n+1} \mathbb{P}\left\{\sum_{i=1}^{n+1} A_{ji} \geq (1-\alpha)(n+1)\right\}$$
by exchangeability
$$= \mathbb{E}\left[\frac{1}{n+1} \sum_{j=1}^{n+1} \mathbb{1}\left\{\sum_{i=1}^{n+1} A_{ji} \geq (1-\alpha)(n+1)\right\}\right]$$

$$= \# \text{ of teams that win } \geq (1-\alpha)(n+1) \text{ games}$$

Step 4: a deterministic bound

How many teams can win $\geq (1 - \alpha)(n+1)$ many games?

Let
$$S = \{j : \sum_{i} A_{ji} \ge (1 - \alpha)(n + 1)\}$$

$$|S| \cdot (1 - \alpha)(n + 1) \le \sum_{j \in S} \sum_{i=1}^{n+1} A_{ji}$$

$$= \sum_{j \in S} \sum_{i \in S} A_{ji} + \sum_{j \in S} \sum_{i \in S^{c}} A_{ji}$$

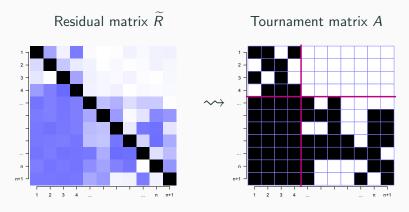
$$= \frac{1}{2} \cdot \sum_{j \in S} \sum_{i \in S} \underbrace{(A_{ji} + A_{ij})}_{\le 1} + \sum_{j \in S} \sum_{i \in S^{c}} A_{ji}$$

$$= \frac{1}{2} \cdot \sum_{j \in S} \sum_{i \in S} \underbrace{(A_{ji} + A_{ij})}_{\le 1} + \sum_{j \in S} \sum_{i \in S^{c}} A_{ji}$$

Jackknife+: intuition for the factor of 2

The worst case scenario, to achieve $|S| \approx 2\alpha(n+1)$:

- ullet pprox 2lpha(n+1) "good" teams, pprox (1-2lpha)(n+1) "bad" teams
- A "good" team always beats a "bad" team
- All other games are random



Jackknife+: intuition for the factor of 2

This worst case scenario is connected to *instability*— $\widehat{\mu} = \mathcal{A}(Z_1, \dots, Z_n)$ is very sensitive to changing a single Z_i

Among "good" teams $i \neq j \neq k$:

• May have
$$\widetilde{R}_{ij} > \widetilde{R}_{ji}$$
 and $\widetilde{R}_{jk} > \widetilde{R}_{kj}$ and $\widetilde{R}_{ki} > \widetilde{R}_{ik}$ and $\widetilde{R}_{ki} > \widetilde{R}_{ik}$ " Y_i is harder to predict than Y_i " predict than Y_i "

We will return to this later.

Cross-validation based methods

CV+ and cross-conformal

From leave-one-out to *K*-fold

To avoid computational cost of leave-one-out CV — K-fold CV (e.g., K=5 or K=10)

- Partition $\{1,\ldots,n\}$ into K folds $A_1\cup\cdots\cup A_K$
- Fit model $\widehat{\mu}_{-A_k} = \mathcal{A}\Big(\{(X_i,Y_i): i\in\{1,\ldots,n\}\backslash A_k\}\Big)$
- For $i \in A_k$ define $R_i = |Y_i \widehat{\mu}_{-A_k}(X_i)|$

$$\mathcal{C}(X_{n+1}) = \widehat{\mu}(X_{n+1}) \pm \mathsf{Quantile}_{1-\alpha}(R_1, \dots, R_n)$$

From leave-one-out to *K*-fold

Generalize jackknife+ to the K-fold setting \rightsquigarrow CV+

K-fold CV+3

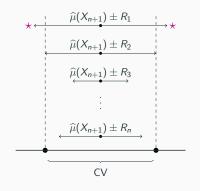
- Partition $\{1,\ldots,n\}$ into K folds $A_1\cup\cdots\cup A_K$
- Fit model $\widehat{\mu}_{-A_k} = \mathcal{A}\Big(\{(X_i,Y_i): i\in\{1,\ldots,n\}\backslash A_k\}\Big)$
- For $i \in A_k$ define $R_i = |Y_i \widehat{\mu}_{-A_k}(X_i)|$
- Prediction set

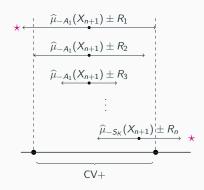
$$\mathcal{C}(X_{n+1}) = \left[-\mathsf{Quantile}_{(1-\alpha)(1+1/n)} \left(\left\{ -\widehat{\mu}_{-A_k}(X_{n+1}) + R_i \right\} \right),$$

$$\mathsf{Quantile}_{(1-\alpha)(1+1/n)} \left(\left\{ \widehat{\mu}_{-A_k}(X_{n+1}) + R_i \right\} \right) \right]$$

³B., Candès, Ramdas, Tibshirani 2019, Predictive inference with the jackknife+

From leave-one-out to *K*-fold





Cross-conformal prediction

CV+ is related to a more general method:

Cross-conformal prediction^{4,5}

- Partition $\{1,\ldots,n\}$ into K folds $A_1\cup\cdots\cup A_K$
- Fit score function $s^{(k)} = \mathcal{A}\Big(\{(X_i,Y_i): i \in \{1,\ldots,n\} \setminus A_k\}\Big)$
- For $i \in A_k$ define $S_i = s^{(k)}(X_i, Y_i)$
- Prediction set

$$C(X_{n+1}) = \left\{ y \in \mathcal{Y} : \sum_{k=1}^K \sum_{i \in A_k} \mathbb{1} \left\{ S_i \ge s^{(k)}(X_{n+1}, y) \right\} \ge \alpha(n+1) \right\}$$

⁴Vovk 2015, Cross-conformal predictors

⁵Vovk et al 2018, Cross-conformal predictive distributions

Cross-conformal prediction

For the residual score function $s(x, y) = |y - \widehat{\mu}(x)|$,

$$C_{\mathsf{cross\text{-}conf.}}(X_{n+1}) \subseteq C_{\mathsf{CV}+}(X_{n+1})$$

Proof (for the case K = n, i.e., jackknife+)

Recall from the proof of the jackknife+ theorem:

$$Y_{n+1} \notin \mathcal{C}_{j+}(X_{n+1}) \Rightarrow \sum_{i=1}^{n} \mathbb{1}\left\{\widetilde{R}_{n+1,i} > \widetilde{R}_{i,n+1}\right\} \geq (1-\alpha)(n+1)$$

Rewrite as:

$$Y_{n+1} \in \mathcal{C}_{j+}(X_{n+1}) \iff \sum_{i=1}^{n} \mathbb{1} \left\{ \widetilde{R}_{n+1,i} > \widetilde{R}_{i,n+1} \right\} \le (1-\alpha)(n+1)-1$$

$$= \sum_{i=1}^{n} \mathbb{1} \left\{ s^{(i)}(X_{n+1}, Y_{n+1}) > S_{i} \right\}$$

$$= n - \sum_{i=1}^{n} \mathbb{1} \left\{ S_{i} \ge s^{(i)}(X_{n+1}, Y_{n+1}) \right\}$$

$$> \alpha(n+1) \iff Y_{n+1} \in \mathcal{C}_{cross-conf}(X_{n+1})$$

Cross-validation based methods

Theory for cross-conformal (and CV+)

Theorem: coverage for CV+ and cross-conformal

If Z_1, \ldots, Z_{n+1} are i.i.d., and A is symmetric, then K-fold cross-conformal satisfies

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}(X_{n+1})\right\} \ge \begin{cases} 1 - 2\alpha - 2/K^{-6} \\ 1 - 2\alpha - 2K/n^{-7} \end{cases}$$

As a special case, the same is true for K-fold CV+.

For any K, then,

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}(X_{n+1})\right\} \ge 1 - 2\alpha - \frac{2}{\sqrt{n}}.$$

⁶B., Candès, Ramdas, Tibshirani 2019, Predictive inference with the jackknife+

⁷Vovk et al 2018, Cross-conformal predictive distributions

Theorem part 1—proved via tournament matrix

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}(X_{n+1})\right\} \ge 1 - 2\alpha - 2/K$$

Imagine that we sample additional test data:

$$\underbrace{(X_1,Y_1),\ldots,(X_n,Y_n),}_{K \text{ training folds } A_1,\ldots,A_K}\underbrace{(X_{n+1},Y_{n+1}),\ldots,(X_{n+n/K},Y_{n+n/K})}_{1 \text{ test fold } A_{K+1}} \overset{\mathrm{iid}}{\sim} P$$

For each $k \neq k' \in \{1, \dots, K+1\}$ define

$$\widetilde{s}_{-kk'} = \mathcal{A}\left((Z_i)_{i \in [n+n/K] \setminus (A_k \cup A_{k'})}\right)$$

and define scores, $\widetilde{S}_{ij} = \widetilde{s}_{-kk'}(X_i, Y_i)$ for $i \in A_k$, $j \in A_{k'}$

For a permutation σ on [n+n/K], we say that σ preserves folds if

$$i,j$$
 in same fold $\iff \sigma(i),\sigma(j)$ in same fold

If σ preserves folds, and Π is the corresponding perm. matrix,

$$\widetilde{S} \stackrel{\mathsf{d}}{=} \Pi \widetilde{S} \Pi^{\top}$$

Define the tournament matrix $A \in \{0,1\}^{(n+n/K)\times(n+n/K)}$:

$$A_{ij} = \mathbb{1}\left\{\widetilde{S}_{ij} > \widetilde{S}_{ji}
ight\}$$
 (or $A_{ij} = 0$ if i,j in same fold)

 $\Rightarrow A \stackrel{d}{=} \Pi A \Pi^{\top}$ if Π preserves folds

Verify via construction of $C(X_{n+1})$:

$$Y_{n+1} \notin \mathcal{C}(X_{n+1}) \iff \sum_{i=1}^{n+n/K} A_{n+1,i} \geq (1-\alpha)(n+1)$$

$$\mathbb{P}\left\{Y_{n+1} \not\in \mathcal{C}(X_{n+1})\right\} = \mathbb{P}\left\{ \begin{aligned} &\text{in a tournament with } n+n/K \\ &\text{teams, team } n+1 \text{ wins} \\ &\geq \underbrace{\left(1-\alpha\right)\!\left(n+1\right)}_{=} \text{ games} \end{aligned} \right\} \\ &= (1-[\alpha+(1-\alpha)\frac{n/K-1}{n+n/K}])(n+n/K)$$

$$\leq 2\left[\alpha + (1-\alpha)\frac{n/K-1}{n+n/K}\right] \leq 2\alpha + 2/K$$

via counting argument (as for jackknife+ proof)

Theorem part 2—proved via averaging p-values

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}(X_{n+1})\right\} \ge 1 - 2\alpha - 2K/n$$

$$C(X_{n+1}) = \left\{ y \in \mathcal{Y} : 1 + \sum_{k=1}^{K} \sum_{i \in A_k} 1 \left\{ S_i \ge s^{(k)}(X_{n+1}, y) \right\} / \nearrow \alpha(n+1) \right\}$$
$$= \left\{ y \in \mathcal{Y} : \frac{1 + \sum_{k=1}^{K} \sum_{i \in A_k} 1 \left\{ S_i \ge s^{(k)}(X_{n+1}, y) \right\}}{n+1} > \alpha \right\}$$

For each fold k = 1, ..., K define a p-value

$$p_{k}^{y} = \frac{1 + \sum_{i \in A_{k}} \mathbb{1}\left\{S_{i} \geq s^{(k)}(X_{n+1}, y)\right\}}{\frac{n}{K} + 1}$$

- As for split CP, $p_k^{Y_{n+1}}$ is a valid p-value (holdout & test scores are exch.)
- Define $\bar{p}^y = \frac{1}{K} \sum_{k=1}^K p_k^y$ Then $\bar{p}^{Y_{n+1}}$ is a valid p-value up to a factor of 2: ⁸ $\mathbb{P}\left\{\bar{p}^{Y_{n+1}} \leq a\right\} \leq 2a, \text{ for all } a \in [0,1].$

⁸Rüschendorf 1982, Random variables with maximum sums; Vovk & Wang 2020, Combining p-values via averaging

$$C(X_{n+1}) = \left\{ y \in \mathcal{Y} : \frac{1 + \sum_{k=1}^{K} \sum_{i \in A_k} \mathbb{1}\left\{S_i \ge s^{(k)}(X_{n+1}, y)\right\}}{n+1} > \alpha \right\}$$

$$= \left\{ y \in \mathcal{Y} : \frac{1}{K} \sum_{k=1}^{K} p_k^y > \alpha + \frac{K-1}{n+K}(1-\alpha) \right\}$$

$$\implies \mathbb{P}\left\{Y_{n+1} \notin \mathcal{C}(X_{n+1})\right\} = \mathbb{P}\left\{\bar{p}^{Y_{n+1}} \le \alpha + \frac{K-1}{n+K}(1-\alpha)\right\}$$
$$\le 2\left(\alpha + \frac{K-1}{n+K}(1-\alpha)\right) \le 2\alpha + 2K/n$$

Cross-validation based methods

Connection to algorithmic stability

Why might jackknife/CV fail to cover? Why might jackknife+/CV+ have coverage $1-2\alpha$ not $1-\alpha$?

Examples we have seen:

- A = least squares for even n, neural net for odd n
- Least squares with $n \approx d$

These examples are unstable—

 $\widehat{\mu} = \mathcal{A}(Z_1, \dots, Z_n)$ is very sensitive to changing a single Z_i

Algorithmic stability⁹

Algorithm \mathcal{A} is (ϵ, ν) -stable if

$$\mathbb{P}\left\{\left|\widehat{\mu}(X_{n+1}) - \widehat{\mu}_{-i}(X_{n+1})\right| \le \epsilon\right\} \ge 1 - \nu$$

Note that this property depends on both ${\cal A}$ and distrib. of data

- Example: ridge regression is stable (due to strong convexity)
- Example: K-nearest-neighbors is stable (if $K \ll n$)
- Next week: stability of bootstrap / ensembled methods

⁹Bousquet & Elisseeff 2002, Stability and Generalization

Define the inflated jackknife interval:

$$\mathcal{C}^{\epsilon}(X_{n+1}) = \widehat{\mu}(X_{n+1}) \pm \left(\mathsf{Quantile}_{1-\alpha}(R_i) + \epsilon\right)$$

Theorem: jackknife under stability¹⁰

If ${\mathcal A}$ satisfies (ϵ, ν) -stability, then

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}^{\epsilon}(X_{n+1})\right\} \ge 1 - \alpha - 2\sqrt{\nu}$$

(similar results hold for jackknife+)

¹⁰B., Candès, Ramdas, Tibshirani 2019, Predictive inference with the jackknife+; see also Steinberger & Leeb, Conditional predictive inference for stable algorithms

A related method:

The conformal jackknife

Consider a leave-one-out version of full conformal:

$$\widetilde{\mathcal{A}}$$
: data $(Z_1,\ldots,Z_{n+1}) \mapsto \text{scores } (S_1,\ldots,S_{n+1})$

where

$$S_i = |Y_i - [\mathcal{A}(Z_1, \dots, Z_{i-1}, Z_{i+1}, \dots, Z_{n+1})](X_i)|$$

If $\ensuremath{\mathcal{A}}$ overfits, this performs better than full CP with residual score Validity:

- ullet Symmetry of ${\mathcal A} \ \Rightarrow \$ symmetry of $\widetilde{{\mathcal A}}$
- \leadsto if data is exch., then $\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}_{\mathsf{conf. jack.}}(X_{n+1})\right\} \geq 1 \alpha_{35/39}$

Proof for jackknife under stability:

Key idea: relate $C_{\mathsf{jack}}^{\epsilon}(X_{n+1})$, to conformal jackknife run at $\alpha' > \alpha$

$$\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}_{\mathsf{jack}}^{\epsilon}(X_{n+1})\right\} \geq \mathbb{P}\left\{Y_{n+1} \notin \mathcal{C}_{\mathsf{jack-CP}}^{\alpha'}(X_{n+1})\right\} \\ - \mathbb{P}\left\{\mathcal{C}_{\mathsf{jack}}^{\epsilon}(X_{n+1}) \not\supseteq \mathcal{C}_{\mathsf{jack-CP}}^{\alpha'}(X_{n+1})\right\}$$

Define models & residuals:

$$\widehat{\mu}_{-i} = \mathcal{A}(Z_1, \dots, Z_{i-1}, Z_{i+1}, \dots, Z_n) \qquad \rightsquigarrow R_i = |Y_i - \widehat{\mu}_{-i}(X_i)|$$

$$\widetilde{\mu}_{-i} = \mathcal{A}(Z_1, \dots, Z_{i-1}, Z_{i+1}, \dots, Z_n, Z_{n+1}) \qquad \rightsquigarrow S_i = |Y_i - \widetilde{\mu}_{-i}(X_i)|$$

The intervals:

$$\begin{split} \mathcal{C}^{\epsilon}_{\mathsf{jack}}(X_{n+1}) &= \widehat{\mu}(X_{n+1}) \pm \mathsf{Quantile}_{1-\alpha}(R_i + \epsilon) \\ \mathcal{C}^{\alpha'}_{\mathsf{jack-CP}}(X_{n+1}) &= \widehat{\mu}(X_{n+1}) \pm \mathsf{Quantile}_{(1-\alpha')(1+1/n)}(S_i) \end{split}$$

$$\Rightarrow \mathcal{C}^{\epsilon}_{\mathsf{jack}}(X_{n+1}) \supseteq \mathcal{C}^{\alpha'}_{\mathsf{jack-CP}}(X_{n+1}) \text{ if } \underbrace{R_i + \epsilon \geq S_i \text{ for sufficiently many } i}_{\mathsf{holds w.h.p. under stability assumption}}$$

Summary

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Summary & preview

Summary: lectures 1, 2, 3

Lecture 1–3 topics:

- The exchangeability framework
- The distribution-free prediction problem
- Versions of the methodology: split CP, full CP, and CV-type conformal methods

Conformal methods allow for distribution-free predictive inference using any base model, & assuming only exchangeability

Preview: lectures 4, 5, 6

All methods thus far have focused on predictive inference with marginal coverage:

$$\underbrace{\mathbb{P}\left\{Y_{n+1} \in \mathcal{C}(X_{n+1})\right\}}_{\text{the goal is always coverage}} \geq \underbrace{1-\alpha}_{\text{the guarantee is always averaged over the distrib.}}_{\text{of training} + \text{ test data}}$$

- Lecture 4: beyond predictive coverage (& more next week)
- Lectures 5 & 6: beyond the marginal coverage guarantee