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# Conformal prediction for surrogate modelling in the UQ framework

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• For a computer code  $g: \mathcal{X} \to \mathcal{Y}$  used in industrial applications, the Uncertainty Quantification (UQ) methodology aims at evaluating how uncertainty on the inputs  $X \in \mathcal{X}$  affects our knowledge of the output  $g(X) \in \mathcal{Y}$  [De Rocquigny et al., 2008].

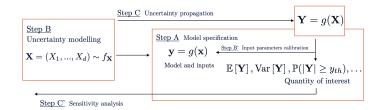


Figure - General UQ methodology.

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- Since these codes are time-costly, surrogates  $\widetilde{g}$  are constructed for performing heavy simulations like Monte-Carlo batch runs for Step C, C'.
- For objective reasons, one needs to assess the quality of these surrogates.
- We propose to use the Conformal prediction paradigm [Vovk et al., 2005] which is a generic, model-agnostic theory allowing to build prediction sets to these surrogates with frequentist coverage guarantees.

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[Vovk et al., 2005]. We fix a probability space  $(\Omega, \mathcal{F}, \mathbb{P})$ .

#### Definition

Let  $\mathcal{X},\mathcal{Y}$  be metric spaces and  $\mathcal{Z}=\mathcal{X}\times\mathcal{Y}$ . Let  $n\in\mathbb{N}$  and  $\mathcal{D}=\{Z_1,\ldots,Z_n\}\in 2^{\mathcal{Z}}$  a training sample. For  $\alpha\in(0,1)$  a conformal predictor of coverage  $\alpha$  is any measurable function of the form :

$$C_{\alpha} \colon 2^{\mathcal{Z}} \times \mathcal{X} \to 2^{\mathcal{Y}}$$

$$(\mathcal{D}, X) \mapsto C_{n,\alpha}(X),$$

$$(1)$$

such that for a new point  $Z_{n+1} = (X_{n+1}, Y_{n+1}) \in \mathcal{Z}$ :

$$\mathbb{P}\left(Y_{n+1} \in C_{n,\alpha}(X_{n+1})\right) \ge 1 - \alpha. \tag{2}$$

Three main methods exist to estimate these set-functions : full-conformal, split-conformal and cross-conformal estimators. We focus on the latter.

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## Full-Conformalized Ridge Regression (CRR)

[Vovk et al., 2005] A non-conformity score is any measurable function of the form :

$$A: 2^{\mathbb{Z}} \times \mathbb{Z} \to \overline{\mathbb{R}}$$

$$(\mathcal{D}, Z) \mapsto A(\mathcal{D}, Z).$$
(3)

Assume  $\mathcal{X} = \mathcal{Y} = \mathbb{R}$  and  $R_{\mathcal{D}}$  a regression rule learned on  $\mathcal{D} = \{Z_1, \dots, Z_n\} = \{(X_1, Y_1), \dots, (X_n, Y_n)\}$ . A straightforward non-conformity score is given by the difference between prediction and actual value, for all  $i = 1, \dots, n$ :

$$A(\mathcal{D}, Z_i) = Y_i - R_{\mathcal{D}}(X_i) = Y_i - \widehat{Y}_i \tag{4}$$

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# Full-Conformalized Ridge Regression (CRR)

For a new input  $X_{n+1} \in \mathcal{X}$  and output  $Y \in \mathcal{Y}$ , we denote by  $\widetilde{\mathcal{D}} := \mathcal{D} \sqcup (X_{n+1}, Y)$ . The full conformal predictor is given for all confidence level  $0 < \alpha < 1$  by :

$$\widehat{C}_{n,\alpha}^{CRR}(X_{n+1}) = \left\{ Y \in \mathcal{Y} : \frac{\#\{i : A(\widetilde{\mathcal{D}}, Z_i) \ge A(\widetilde{\mathcal{D}}, (X_{n+1}, Y))\}}{n} > \alpha \right\}.$$

- This estimator is based on the "dissimilarity" of the new pair inside the updated dataset. It achieves the required coverage property.
- It is computationally intractable due to the full grid search on the label space  $\mathcal{Y}$ .
- To tackle this, more advanced estimators are proposed in the following.
- In the rest of this talk we suppose that Y = g(X) where g is some deterministic function (e.g a numerical simulation code).

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[Barber et al., 2021]. We train a metamodel  $\widehat{g}$  on  $\mathcal{D}$  and n leave-one-out metamodels  $\widehat{g}_{-i}$  on  $\mathcal{D}\setminus(X_i,g(X_i))$ . The Jackknife+ estimator is given by :

$$\widehat{C}_{n,\alpha}^{J+}(X_{n+1}) = \left[ q_{n,\alpha}^{-} \left\{ \widehat{g}_{-i}(X_{n+1}) - R_{i}^{LOO} \right\}, q_{n,\alpha}^{+} \left\{ \widehat{g}_{-i}(X_{n+1}) + R_{i}^{LOO} \right\} \right]$$
(5)

Where the leave-one-out error defined by :

$$R_i^{LOO} := |g(X_i) - \widehat{g}_{-i}(X_i)| \tag{6}$$

- With this estimator we have a coverage guarantee of  $1-2\alpha$ .
- However, these intervals have almost constant width for any new point.

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- Let  $K \in \{1, ..., n\}$ . We divide  $\mathcal{D}^{\text{train}}$  in K-disjoint subsets  $S_1 \cup ... \cup S_K$ .
- We fit K metamodels with the k-th fold removed :  $\widehat{g}_{-S_k}$ .
- We compute the conformity scores :

$$R_i^{CV} := |g(X_i) - \widehat{g}_{-S_{k(i)}}(X_i)|, \ \forall i \in \{1, \dots, n\},$$
 (7)

where  $S_{k(i)}$  is the fold containing  $X_i$ .

• We estimate our prediction intervals :

$$\widehat{C}_{n,\alpha}^{\mathsf{CV+}}(X_{n+1}) = \left[\widehat{q}_{n,\alpha}^{\pm} \left\{ \widehat{g}_{-S_{k(i)}}(X_{n+1}) \pm R_i^{\mathsf{CV}} \right\} \right] \tag{8}$$

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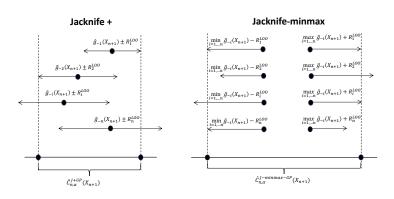
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$$with \, R_i^{LOO} = |Y_i - \hat{g}_{-i}(X_i)|$$

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Additionally, we can add minmax on both the J+ and the CV+ methods by replacing the metamodel prediction with :

$$\widehat{g}_{-i}(X_{n+1}) \longleftrightarrow \min_{i} \widehat{g}_{-i}(X_{n+1}), \max_{i} \widehat{g}_{-i}(X_{n+1}).$$
 (9)

$$\widehat{g}_{-S_k(i)}(X_{n+1}) \longleftrightarrow \min_{i} \widehat{g}_{-S_k(i)}(X_{n+1}), \max_{i} \widehat{g}_{-S_k(i)}(X_{n+1}). \tag{10}$$

The intervals will not be centered anymore and we have the coverage guarantee [Barber et al., 2021] :

$$\mathbb{P}\left(g(X_{n+1}) \in \widehat{C}_{n,\alpha}^{*+-\mathsf{mm}}(X_{n+1})\right) \ge 1 - \alpha. \tag{11}$$

Moreover, the resulting intervals will be more conservative.

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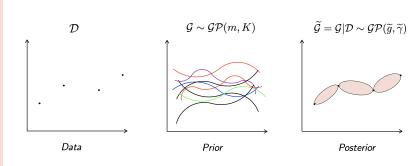
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## Bayesian credibility intervals

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Since we are in a Gaussian setting, we can have access to the confidence intervals for any new point  $X_{n+1} \in \mathcal{X} \setminus X$ :

$$CR_{\alpha}(X_{n+1}) = \left[\widetilde{g}(X_{n+1}) \pm F^{-1}\left(1 - \frac{\alpha}{2}\right)\widetilde{\gamma}(X_{n+1})\right]$$
(12)

which in our case are  $\underline{\text{credibility}}$  intervals. Here F is the CDF of the standard normal distribution.

If g was *truly* drawn from our posterior  $\mathcal{G}|(X, g(X))$ , then we would have the exact coverage :

$$\mathbb{P}\left(g(X_{n+1}) \in \mathcal{CR}_{\alpha}(X_{n+1})\right) = 1 - \alpha,\tag{13}$$

and  $CR_{\alpha}(X_{n+1})$  would be the true prediction interval for the true g.

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In practice we don't have access to the true distribution  $\mathbb{P}_{(X,g(X))}$ . If we dispose of a test dataset :

$$\mathcal{D}^{test} = \{ (X_1, g(X_1)), \dots, (X_m, g(X_m)) \}, \tag{14}$$

different from  $\mathcal{D}$ , then in all generality we cannot expect to have :

$$\frac{1}{m}\sum_{i=1}^{m}1\left\{g(X_{i})\in\mathcal{CR}_{\alpha}(X_{i})\right\}\geq1-\alpha, \text{ a.s.}$$
 (15)

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## The Burnaev-Wasserman program

[Burnaev and Vovk, 2014] Assume that  $\mathcal{X} \subset \mathbb{R}^d$ , for all i,  $X_i \in L^2(\Omega)$  and the model g is truly gaussian. The credibility sets have exact coverage and output an interval of the form :

$$\mathcal{CR}_{\alpha}(X_{n+1}) = [B_*, B^*] \tag{16}$$

The CRR method with the GPR rule outputs a prediction interval of the form :

$$\widehat{C}_{n,\alpha}^{CRR}(X_{n+1}) = [C_*, C^*].$$
 (17)

A natural question is to compare the differences of the bounds of these two intervals and their asymptotic behaviour

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See [Burnaev and Vovk, 2014] for a proof of the following.

#### **Theorem**

Under the above assumptions we get

$$\sqrt{n}\left(B^* - C^*\right) \xrightarrow[n \to \infty]{\text{Law}} \mathcal{N}\left(0, h(\alpha)\right), \tag{18}$$

and similarly for the lower-bound.

Here h is a function of the  $1-\alpha/2$ -quantile of the standard normal distribution and of the mean and variance of the input distribution.

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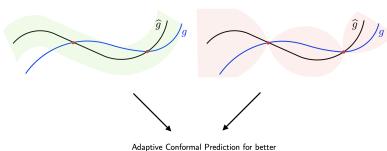
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#### Gaussian Processes



$$\mathbb{P}\left(g(X) \in C_{\widehat{g},\alpha}(X)\right) \ge 1 - \alpha. \tag{19}$$

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The idea is to adapt the Jackknife+ method presented earlier to GPR metamodels to have adaptive prediction intervals. We denote these predictors as  $\widehat{C}_{n,\alpha}^{J+GP}$  and can prove the following theorem :

#### **Theorem**

Assume  $\mathcal{D} = (\mathbf{X}, g(\mathbf{X}))$  is exchangeable. For a new point  $X_{n+1} \in \mathcal{X} \setminus \mathbf{X}$  and a coverage level  $\alpha \in (0,1)$ :

$$\mathbb{P}\left(g(X_{n+1}) \in \widehat{C}_{n,\alpha}^{J+GP}(X_{n+1})\right) \ge 1 - 2\alpha. \tag{20}$$

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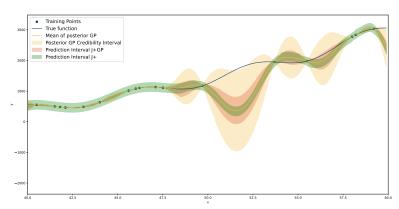
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## Example on a toy function

• The Vincent Blot $^{\bigcirc}$  highly original  $g_{VB}$ :

$$g_{VB}(x) = 3x\sin(x) - 2x\cos(x) + \frac{x^3}{40} - \frac{1}{2}x^2 - 10x,$$
 (21)

which we sample at ease.



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- Conformal prediction is a method for performing distribution-free uncertainty quantification of machine learning algorithms.
- It can be used in a Bayesian regression setting for testing the soft-assumption on the original model.
- Prediction intervals for Gaussian Processes using an adaptation of the CP algorithms can be built (upcoming paper [Jaber and Blot, 2023] - in progress).
- A more robust uncertainty quantification of this type of surrogates can be achieved and can thus serve in assessing their quality for the purpose of performing industrial UQ.

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