Uncertainty Quantification in Machine Learning From Aleatoric to Epistemic

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- Learning in the sense of generalising beyond the data seen so far is necessarily based on a process of induction.
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- Other sources of uncertainty exist: incorrect model assumptions (model misspecification), noisy or imprecise data, etc.
- Trustworthy representation of uncertainty is desirable and should be considered as a key feature of any machine learning method, all the more in safety-critical application domains.

Self-awareness of ML systems

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- Many applications require safe and reliable predictions, and hence a certain level of self-awareness of ML systems:
 - equip predictions with an appropriate quantification of uncertainty,
 - reject a decision in cases of high uncertainty (abstention) ,
 - deliver a credible set-valued prediction (partial abstention),

▶ ...



Driver assistance systems: a safety-critical application

Adversarial examples

There is really but one thing to say about **this** sorry movie It should never have been made The first one one of my favourites An American Werewolf in London is a great movie with a good plot good actors and good FX But this one It stinks to heaven with a cry of helplessness

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The adversarial example (right) is misclassified by a machine learning model trained on textual data, which changes its prediction due to a change of a single (actually unimportant) word (Sato *et al.*, 2018).

Lack of uncertainty-awareness

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Predictions by EfficientNet on test images from ImageNet: For the left image, the neural network predicts "typewriter keyboard" with certainty 83.14%, for the right image "stone wall" with certainty 87.63%.





Levels of self-awareness and uncertainty representation



Probability distributions p = (p(a), p(b), p(c)) on $\Omega = \{a, b, c\}$, for example $\Omega = \{$ home wins, draw, away wins $\}$, as points in a Barycentric coordinate system.

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Assuming a probabilistic data generating process $p(\mathbf{x}, y) = p(\mathbf{x})p(y | \mathbf{x})$, probabilistic predictors (estimating $p(y | \mathbf{x})$) are natural primitives.



A learner is given access to a set of (i.i.d.) training data

$$\mathcal{D} := \left\{ (\pmb{x}_1, y_1), \ldots, (\pmb{x}_N, y_N)
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where \mathcal{X} is an instance space and \mathcal{Y} the set of outcomes. Given a hypothesis space $\mathcal{H} \subset \mathcal{Y}^{\mathcal{X}}$ and a loss function

$$\ell: \mathcal{Y} \times \mathcal{Y} \longrightarrow \mathbb{R},$$

the goal of the learner is to induce a hypothesis $h^* \in \mathcal{H}$ with low **risk**

$$R(h) := \int_{\mathcal{X} \times \mathcal{Y}} \ell(h(\mathbf{x}), y) \, d \, P(\mathbf{x}, y) \, .$$

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If predictions are probabilities

$$h(oldsymbol{x}) = \hat{oldsymbol{
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the loss is defined on $\Delta_K \times \mathcal{Y}$; a common example is the logistic loss (log-loss)

$$\ell(\hat{\boldsymbol{p}}, y) = -\log(p_y).$$

The learner's choice of a hypothesis is commonly guided by the empirical risk

$$R_{emp}(h) := rac{1}{N} \sum_{i=1}^{N} \ell(h(oldsymbol{x}_i), y_i) \; .$$

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Yet, since R_{emp}(h) is only an estimation of the true risk R(h), the (regularised) empirical risk minimiser

$$\hat{h} := rgmin_{h \in \mathcal{H}} R_{emp}(h) + \Omega(h)$$

will normally not coincide with the true risk minimiser

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Correspondingly, there remains **uncertainty** regarding h^* , the approximation quality of \hat{h} (in the sense of its proximity to h^*) and its true risk $R(\hat{h})$, as well as **predictive uncertainty** about \hat{y}_q for an individual **query instance** $\mathbf{x}_q \in \mathcal{X}$.

A precise specification of the problem setting and underlying assumptions is an important prerequisite, not only for providing learning guarantees, but also for uncertainty quantification.



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- Here, one might be quite sure about the class of the query under standard assumptions of binary classification, but much less so in a setting of **novelty detection**, where new classes may emerge.
- Likewise, assumptions such as i.i.d. data generation are really crucial (the past should be representative of the future).

Sources of uncertainty
• A query instance \mathbf{x}_q gives rise to a conditional probability on \mathcal{Y} :

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- Thus, even given full information in the form of the measure P (and its density p), uncertainty about the actual outcome y remains.
- The best point predictions (minimizing expected loss) are prescribed by the pointwise Bayes predictor f*:

$$f^*(\mathbf{x}) := \operatorname*{arg\,min}_{\hat{y} \in \mathcal{Y}} \int_{\mathcal{Y}} \ell(y, \hat{y}) \, dP(y \mid \mathbf{x}) \, .$$

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- We shall refer to this uncertainty as **model uncertainty**.

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- We shall refer to this uncertainty as model uncertainty.
- Due to model uncertainty, one cannot guarantee

 $h^*(\boldsymbol{x})=f^*(\boldsymbol{x}),$

or, in the case of probabilistic predictions $p^*(y \mid \boldsymbol{x}) := p(y \mid \boldsymbol{x}, h^*)$, that

 $p^*(\cdot \mid \boldsymbol{x}) = p(\cdot \mid \boldsymbol{x}).$

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• We refer to the uncertainty about the discrepancy between \hat{h} and h^* as approximation uncertainty.





Agenda

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- 2. Training probabilistic predictors
- 3. Calibration
- 4. Set-valued (conformal) prediction
- 5. Epistemic uncertainty

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- Captures stochastic nature of dependence p(y | x) between instances and outcomes, and hence aleatoric uncertainty.
- Yet, pretends full certainty about this dependence, thereby ignoring approximation and model uncertainty (epistemic uncertainty).

Suppose that hypotheses *h* are (uniquely) identified by parameters $\theta \in \Theta$, i.e.,

$$\mathcal{H} = \{h_{\boldsymbol{ heta}} \,|\, \boldsymbol{ heta} \in \Theta\}$$
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Learning (model induction) then comes down to parameter estimation.

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The maximum likelihood (ML) principle suggests to pick the parameter with the highest likelihood:

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- In general, ML estimation has good statistical properties.
- Under the i.i.d. assumption, ML estimation comes down to solving

$$\hat{\boldsymbol{\theta}} = \arg\max_{\boldsymbol{\theta} \in \Theta} \prod_{i=1}^{N} P((\boldsymbol{x}_i, y_i) | \boldsymbol{\theta}) = \arg\max_{\boldsymbol{\theta} \in \Theta} \sum_{i=1}^{N} \log P((\boldsymbol{x}_i, y_i) | \boldsymbol{\theta})$$

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We can think of the above model as follows: First, an instance x is assigned a (latent) score (expressing the propensity for the positive class)

$$s = s(\mathbf{x}) = \langle \mathbf{ heta}, \mathbf{x}
angle,$$

which is then transformed into a probability via the (logistic) link function

$$\phi(s) = \frac{1}{1 + \exp(-s)}$$

The probability of a label y_i depends on the distance of x_i from the hyperplane defined by $\theta^{\top} x = 0$.



In order to learn θ , we can invoke the ML principle:

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Probability is only maximised over the y_n, while the x_n are assumed to be fixed.
 Logistic regression is an example of discriminative learning, i.e., it learns a map

$$h: \mathbf{x} \mapsto p(Y \mid \mathbf{x}).$$

This is to be distinguished from **generative** learning, which essentially means learning the entire data-generating process in the form of the joint distribution p(X, Y).

Bayesian learning
■ In the Bayesian approach, learning corresponds to turning a **prior distribution** on *H* into a **posterior**:

$$p(h \mid \mathcal{D}) = \frac{p(\mathcal{D} \mid h) p(h)}{p(\mathcal{D})} \propto p(\mathcal{D} \mid h) p(h)$$

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The predictive posterior distribution on \mathcal{Y} is obtained via **model averaging**:

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- Bayesian inference is very costly but can be done approximately, for example, using ensemble methods.
- An alternative is to commit to the hypothesis with maximum a-posteriori probability (MAP inference):

$$h_{MAP} = rgmax_{h \in \mathcal{H}} p(h \,|\, \mathcal{D})$$
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- We also encode a realisation y of Y in terms of a vector \mathbf{y} with entry 1 on position y = k (i.e., the k^{th} outcome was observed), and all other entries 0.
- Expressed in terms of predicted probability, the log-loss can then also be written as

$$\ell(\hat{\boldsymbol{p}}, y) = -\log \hat{p}_y$$
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■ A loss *l* is a proper scoring rule if the expected loss minimiser coincides with the true probability *p*:

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- Thus, given a query \boldsymbol{x} , a learner penalised by a (strictly) proper scoring rule has an incentive to predict the true (conditional) probability $\boldsymbol{p} = p(Y | \boldsymbol{x})$.
- **•** For example, the **Brier score** $\sum_k (\hat{p}_k y_k)^2$ is strictly proper, because

$$\sum_{k=1}^{K} p_k \cdot \left[(1 - \hat{p}_k)^2 + \sum_{j \neq k} (\hat{p}_j)^2 \right] = \sum_k (\hat{p}_k)^2 + \sum_k p_k (1 - 2\hat{p}_k)^2$$

is minimised by $\hat{p}_k = p_k$ for all $k \in [K]$.

Define the scoring function on probability vectors $\hat{\boldsymbol{p}}, \boldsymbol{p}$ as

$$S(\hat{\boldsymbol{p}}, \boldsymbol{p}) := \mathbb{E}_{\boldsymbol{Y} \sim \boldsymbol{p}} S(\hat{\boldsymbol{p}}, \boldsymbol{Y}) = \sum_{k} S(\hat{\boldsymbol{p}}, \boldsymbol{y}) p_{k},$$

i.e., as the expected score (under ground-truth \boldsymbol{p}).

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is nonnegative, and strictly proper if $d(\hat{\boldsymbol{p}}, \boldsymbol{p}) = 0$ implies $\hat{\boldsymbol{p}} = \boldsymbol{p}$. $\boldsymbol{e}(\boldsymbol{p}) := S(\boldsymbol{p}, \boldsymbol{p})$ is also called **entropy**.

■ For the **log-loss**, we obtain a decomposition into KL-divergence and information (Shannon) entropy:

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The 0/1 loss is a proper but not a strictly proper scoring rule.

■ The expectation of a strictly proper score can be decomposed as follows:

$$\mathbb{E}_{Y \sim \boldsymbol{\rho}} S(\hat{\boldsymbol{\rho}}, Y) = S(\hat{\boldsymbol{\rho}}, \boldsymbol{\rho}) = d(\hat{\boldsymbol{\rho}}, \boldsymbol{\rho}) + S(\boldsymbol{\rho}, \boldsymbol{\rho}),$$

where the entropy $S(\boldsymbol{p}, \boldsymbol{p})$ is the unavoidable part of the loss (due to the need to predict the realisation of a random variable) and $d(\hat{\boldsymbol{p}}, \boldsymbol{p})$ the extra loss.

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Another decomposition is

$$\mathbb{E} S(\hat{\boldsymbol{p}}, Y) = \mathbb{E} d(\hat{\boldsymbol{p}}, \boldsymbol{c}) + \mathbb{E} e(\boldsymbol{c}),$$

where the expectation is taken over $\mathcal{X} \times \mathcal{Y}$, and $c_k := p(y = k | \hat{p})$, i.e., c is the true class distribution on those instances that receive the same prediction \hat{p} .

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E $d(\hat{p}, c)$ is called calibration loss and $\mathbb{E} e(c)$ refinement loss.

Many ML methods naturally yield predictions in the form of scores

 $s_1,\ldots,s_K\in\mathbb{S}\subseteq\mathbb{R},$

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Yet, calibration is a prerequisite for uncertainty-awareness and important for prediction, decision-making, cost-sensitive classification, etc.

Probability estimation with decision trees



Consider the binary case (K = 2) with probabilistic predictions

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- Broadly speaking, averaged over all instances \mathbf{x} for which the learner predicts $\hat{p}_+ = h(\mathbf{x}) = \alpha$, the fraction of positives is indeed α .
- In other words, a predicted probability vector is supposed to match empirical frequencies, at least in the long run.





Examples of miscalibration: bias toward 1/2 (left), systematic underestimation (right).

Reliability diagram



Reliability diagram



As ground-truth probabilities are not observed, binning is needed in practice.

Reliability diagram



As ground-truth probabilities are not observed, binning is needed in practice.
There is a trade-off in the choice of the width of the bins (more data per bin vs. more fine-granular assessment).

Different post-processing (post-hoc) methods have been proposed for the purpose of calibration, i.e., to construct a calibration function

$$C : \mathbb{S} \longrightarrow [0,1],$$

such that $\hat{p}_+ = C(s)$ is a well-calibrated probability estimate for instances x assigned score s = h(x).

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- For learning *C*, a set of **calibration data** is used:

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This data should be different from the training data (used to learn the scoring classifier h). Otherwise, there is a risk of introducing a bias.

Binning offers a first obvious approach: Partition S into bins (intervals) B_1, \ldots, B_M , and define $C(s) = \hat{p}_{J(s)}$, where J(s) denotes the index of the bin of s (i.e., $s \in B_{J(s)}$).

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- **a** $\hat{p}_1, \ldots, \hat{p}_M$ are chosen so as to minimise the **estimated calibration error** (ECE)

$$\mathsf{ECE} = \sum_{m=1}^{M} rac{|B_m|}{N} |\hat{p}_m - ar{p}_m|,$$

where

$$\bar{p}_m = \frac{\sum_{n=1}^{N} [\![s_n \in B_m]\!] [\![y_n = +1]\!]}{\sum_{n=1}^{N} [\![s_n \in B_m]\!]}$$

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Binning is nonparametric and hence flexible, easy to train, and can directly minimise calibration error, albeit at the cost of (increased) groping loss.

Another method is **Platt scaling**, which essentially applies logistic regression to predicted scores $s \in \mathbb{R}$, i.e., it fits a calibration function C such that

$$\mathcal{C}_{lpha,eta}(s) = rac{1}{1+\exp(-lpha\cdot s -eta)}\,,$$

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Platt scaling is fast and easy to implement, but restricted to sigmoidal calibration functions (pushing scores from the center toward the extremes, hence coming with a risk of over-confidence).



Beta calibration

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again minimising log-loss on \mathcal{D}_{cal} .

Although still restricted in a parametric way, Beta calibration is more flexible than Platt scaling and includes inverse sigmoids and the identity map (which helps prevent over-calibration and unnecessary adjustments).

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- Isotonic regression minimises

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- Note that *C* is evaluated only at a finite number of points; in-between, one may (linearly) interpolate or assume a piecewise constant function.
- Isotonic regression is more expensive in terms of training time and memory consumption.

Let the scores observed for calibration be sorted (and without ties), such that

 $s_1 < s_2 < \ldots < s_N$.

We then seek values $c_1 \leq c_2 \leq \ldots \leq c_N$ which minimize

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Initialise one block B_n for each observation (s_n, y_n) ; the value of the block is $c(B_n) = y_n$ and the width is $w(B_n) = 1$.

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- **Initialise** one block B_n for each observation (s_n, y_n) ; the value of the block is $c(B_n) = y_n$ and the width is $w(B_n) = 1$.
- A merge operation combines two blocks B' and B'' into a new block B with width w(B) = w(B') + w(B'') and value

$$c = rac{w(B')c(B') + w(B'')c(B'')}{w(B') + w(B'')}\,.$$

- PAVA iterates the following steps (the description is somewhat simplified to avoid notational overload):
 - (1) Find the first violating pair, namely, adjacent blocks B_i and B_{i+1} such that $c_i > c_{i+1}$; if there is no such pair, then stop.
 - (2) Merge B_i and B_{i+1} into a new block B.
 - (3) If $c(B) < c(B_{i-1})$ for the left neighbor block B_{i-1} , merge also these blocks and continue doing so until no more violations are encountered.
 - (4) Continue with (1).


Pair-adjacent violators algorithm (PAVA)

I Note that, in the case of binary classification, the target values are all in $\{0, 1\}$:



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- Then, however, the problem becomes conceptually more difficult (and is still a topic of ongoing research).
- Some concepts do not immediately generalise, for example isotonic regression (which assumes a ranking on scores, and rankings are inherently bipartite).
- While essentially coinciding for binary classification, the following definitions of calibration (leading to increasingly difficult problems) can be distinguished for more than two classes:
 - Confidence calibration: Calibration of the highest predicted probability
 - Class-wise calibration: Calibration of the marginal probabilities
 - Multi-class calibration: Calibration of the entire vector of predicted probabilities

Agenda

1. Introduction

- 2. Training probabilistic predictors
- 3. Calibration
- 4. Set-valued (conformal) prediction
- 5. Epistemic uncertainty

Conformal prediction (Balasubramanian *et al.*, 2014) is a framework for reliable prediction that is rooted in classical frequentist statistics and hypothesis testing.

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- Instead of point predictions, CP makes set-valued predictions covering the true outcome with high probability.



$$\longrightarrow \qquad P(y \in \{2,3,9\}) \ge 0.9$$

Given a sequence of training observations

$$(x_1, y_1), (x_2, y_2), \dots, (x_N, y_N), (x_{N+1}, \bullet)$$

and a new query \mathbf{x}_{N+1} with unknown outcome y_{N+1} , • is hypothetically replaced by each candidate, i.e., the hypothesis $y_{N+1} = y$ is tested for all $y \in \mathcal{Y}$:

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■ Only those outcomes *y* for which this **hypothesis can be rejected** at a predefined level of confidence are excluded, while those for which the hypothesis cannot be rejected are collected to form the prediction set or **prediction region** $Y \subseteq \mathcal{Y}$.

In conformal prediction, the "strangeness" of a pattern (x_{N+1}, y) is captured in terms of a **nonconformity score**.

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The higher the score, the more "strange" the pattern (x, y), i.e., the less the data point (x, y) conforms to what one would expect to observe.

Example of a nonconformity score based on nearest neighbors:

$$f(\mathbf{x}, y) = \frac{\sum_{i=1}^{k} d_i^+}{\sum_{i=1}^{k} d_i^-},$$

where d_i^+ is the distance from the i^{th} nearest neighbor labeled y, and d_i^- the distance from the i^{th} nearest neighbor labeled differently.



Applying this function to the sequence of observations, with a specific (though hypothetical) choice of $y = y_{N+1}$, yields a sequence of scores

 $\alpha_1, \alpha_2, \ldots, \alpha_N, \alpha_{N+1},$

where $\alpha_i = f(\mathbf{x}_i, y_i)$.

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■ Denote by σ the permutation of $\{1, ..., N + 1\}$ that sorts the scores in increasing order, i.e., such that

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■ Under the assumption that the hypothetical choice of y_{N+1} is in agreement with the true data-generating process, and that this process has the property of **exchangeability**, every permutation σ has the same probability of occurrence.

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- Thus, the hypothesis $y_{N+1} = y$ can be rejected for those candidates y for which $p(y) < \epsilon$.

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- The error bounds are valid by construction, regardless of the nonconformity function.
- However, the choice of this function has an important influence on the efficiency of conformal prediction, that is, the size of prediction regions: The more suitably the nonconformity function f is chosen, the smaller these sets will be.

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- The above validity property is also called **marginal coverage**. The randomisation is over the entire data generation and prediction procedure; thus, the coverage of 1ϵ is neither guaranteed
 - ▶ for a fixed sequence on previous data (coverage can be higher or lower),
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- There are various other extensions of CP, also for conditional coverage. Besides, instead of controlling coverage, there are variants for controlling more general notions of risk (Angelopoulos *et al.*, 2021).
- Uncertainty quantification with conformal prediction is
 - agnostic to the underlying model,
 - agnostic to the underlying data distribution (i.e., distribution-free),
 - valid for the finite sample case (not only asymptotically).

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- **Epistemic** (aka systematic) uncertainty refers to uncertainty caused by a **lack of knowledge**, i.e., to the epistemic state of the agent.

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- **Epistemic** (aka systematic) uncertainty refers to uncertainty caused by a **lack of knowledge**, i.e., to the epistemic state of the agent.
- As opposed to aleatoric uncertainty, epistemic uncertainty can in principle be reduced on the basis of additional information.



"Not knowing the chance of mutually exclusive events and knowing the chance to be equal are two quite different states of knowledge"

Ronald Fisher (1890-1962)



Both types of uncertainty also play an important role in ML, where the learner's state of knowledge strongly depends on the amount of data seen so far ...



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■ ... but also on the underlying model assumptions:



Uncertainty representation: How should the learner represent its (model, predictive) uncertainty, i.e., which mathematical formalisms should be used?

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- Uncertainty quantification: How to quantify the learner's uncertainty in terms of numbers? How to measure and disentangle the different types of uncertainty (aleatoric, epistemic, total)?

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- Predict the next number: 116, 304, 194, 341, 224, 654, 609, 625, 533, 91, 205, 35, 527, 611, 128, 235, 348, 912, 582, 52, 672, 20, 856, 904, 628, 273, 615, 105, 610, 862, 384, 705, 73, 794, 775, 156, ??

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 $x \leftarrow x imes 237 \mod 971$

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- **E** Fixed weights θ lead to a fixed probability $p(\cdot | \mathbf{x}, \theta)$.



The Bayesian approach

- A Bayesian learner maintains a probability distribution over the hypothesis space (probabilistic predictors).
- The less concentrated that distribution, the higher the learner's epistemic uncertainty.



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Posterior predictive distribution



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■ What we seek is a **decomposition**



One idea is to model epistemic uncertainty as mutual information between outcomes and hypotheses (Depeweg et al., 2018):

H[Y] $= I(Y;\Theta) + H[Y | \Theta]$

epistemic

total uncertainty

aleatoric

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- Intuitively, epistemic uncertainty thus captures the amount of information about the model parameters θ that would be gained through knowledge of the true outcome y.
- Total uncertainty = entropy of the predictive posterior distribution, in the case of discrete *Y* given by

$$\mathsf{TU}(\mathbf{x}) = H[p(y | \mathbf{x})] = -\sum_{y \in \mathcal{Y}} p(y | \mathbf{x}) \log_2 p(y | \mathbf{x}).$$

This uncertainty also includes the (epistemic) uncertainty about the network weights θ , but fixing a set of weights, i.e., considering a distribution $p(y | x, \theta)$, removes the epistemic uncertainty.

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- Therefore, the expectation over the entropies of these distributions,

$$\mathbb{E}_{p(\boldsymbol{\theta} \mid \mathcal{D})} H[p(y \mid \boldsymbol{x}, \boldsymbol{\theta})] = \\ = -\int p(\boldsymbol{\theta} \mid \mathcal{D}) \left(\sum_{y \in \mathcal{Y}} p(y \mid \boldsymbol{x}, \boldsymbol{\theta}) \log_2 p(y \mid \boldsymbol{x}, \boldsymbol{\theta}) \right) d \boldsymbol{\theta} ,$$

is a measure of the aleatoric uncertainty (conditional entropy).

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- Therefore, the expectation over the entropies of these distributions,

$$p(\theta \mid D) H[p(y \mid \mathbf{x}, \theta)] =$$

$$= -\int p(\theta \mid D) \left(\sum_{y \in \mathcal{Y}} p(y \mid \mathbf{x}, \theta) \log_2 p(y \mid \mathbf{x}, \theta) \right) d\theta ,$$

is a measure of the aleatoric uncertainty (conditional entropy).Finally, the epistemic uncertainty is obtained as the difference

$$\mathsf{EU}(\mathbf{x}) := H\big[p(y \,|\, \mathbf{x})\big] - \mathbb{E}_{p(\boldsymbol{\theta} \,|\, \mathcal{D})} H\big[p(y \,|\, \mathbf{x}, \boldsymbol{\theta})\big]$$

which equals the **mutual information** between y and θ .

Example: coin flipping

■ Tossing a coin with bias *p*, task is to predict the next outcome, hypothesis space equipped with Dirichlet distribution



Remarks

- Is a uniform distribution on $\Delta = \mathbb{P}(\mathcal{Y})$ (set of all distributions *p* on \mathcal{Y}) an adequate representation of **complete ignorance** (full epistemic uncertainty)?
- Averaging the (conditional) entropies over all p is meaningful only if all p are indeed known to be equally likely (aleatoric uncertainty is always 1/2).
- **B** But this is certainly not the case, as only one p^* can be the ground truth.



Remarks

- One may also question the **additive decomposition** TU = AU + EU itself.
- In the beginning, total uncertainty should be full (TU = 1), and so should epistemic uncertainty (EU = 1) but this implies AU = 0.
- This suggests a role of AU as a **lower bound** on (the true) aleatoric uncertainty.
- Indeed, epistemic uncertainty partially comprises aleatoric uncertainty (high EU implies high uncertainty about AU, showing interaction between both).

Ensemble methods for uncertainty quantification



Ensemble methods for uncertainty quantification



Ensemble can be seen as an approximation of a distribution.

Ensemble methods for uncertainty quantification



Ensemble can be seen as an approximation of a distribution.Intuitively, diversity is an indicator of epistemic uncertainty.

Based on an ensemble of hypotheses h_1, \ldots, h_M , producing respective predictions p_1, \ldots, p_M , an approximation of conditional entropy can be obtained by

$$\mathsf{AU}(\boldsymbol{x}) := -\frac{1}{M} \sum_{i=1}^{M} \sum_{y \in \mathcal{Y}} p_i(y \mid \boldsymbol{x}) \log_2 p_i(y \mid \boldsymbol{x}),$$

an approximation of total uncertainty (Shannon entropy) by

$$\mathsf{TU}(\boldsymbol{x}) := -\sum_{y \in \mathcal{Y}} \underbrace{\left(\frac{1}{M} \sum_{i=1}^{M} p_i(y \mid \boldsymbol{x})\right)}_{\bar{p}(y \mid \boldsymbol{x})} \log_2 \underbrace{\left(\frac{1}{M} \sum_{i=1}^{M} p_i(y \mid \boldsymbol{x})\right)}_{\bar{p}(y \mid \boldsymbol{x})},$$

and an approximation of epistemic uncertainty (mutual information) by the difference.

Epistemic uncertainty thus defined is equivalent to Jensen-Shannon divergence of the distributions $p_i(y | \mathbf{x})$, i = 1, ..., M



■ For neural networks, it has been shown that techniques such as **Dropout** (Gal and Ghahramani, 2016) and **DropConnect** (Mobiny *et al.*, 2021) can be interpreted as (implicit) ensemble methods, and can hence be used to implement this approach.

- For neural networks, it has been shown that techniques such as **Dropout** (Gal and Ghahramani, 2016) and **DropConnect** (Mobiny *et al.*, 2021) can be interpreted as (implicit) ensemble methods, and can hence be used to implement this approach.
- Of course, any other ensemble technique could be used as well.



We proposed an implementation based on Random Forests, using decision trees that predict probabilities in terms of (Laplace-corrected) relative frequencies (Shaker and Hüllermeier, 2020).



We proposed an implementation based on Random Forests, using decision trees that predict probabilities in terms of (Laplace-corrected) relative frequencies (Shaker and Hüllermeier, 2020).



Empirically, there are no significant performance differences between neural networks and random forests.

Consider a (level-1) loss for probabilistic predictions:

 $\ell_1: \mathbb{P}(\mathcal{Y}) imes \mathcal{Y} \longrightarrow \mathbb{R}$

- Recall that ERM yields good (unbiased) predictors if ℓ_1 is a (strictly) proper scoring rule, which incentivises the learner to predict the true p(y | x).
- **Question**: Can we do the same on the **epistemic level**, i.e., training a predictor

$$h: \mathcal{X} \longrightarrow \mathbb{P}(\mathbb{P}(\mathcal{Y}))$$

by minimising a level-2 loss

$$\ell_2: \mathbb{P}(\mathbb{P}(\mathcal{Y})) \times \mathcal{Y} \longrightarrow \mathbb{R},$$

such that the predictor represents its epistemic uncertainty in a faithful way?

■ A Dirichlet distribution Dir(α) is specified by means of $K \ge 2$ positive real-valued parameters, i.e., a vector $\alpha = (\alpha_1, \ldots, \alpha_K) \in \mathbb{R}_+^K$.

- A Dirichlet distribution Dir(α) is specified by means of K ≥ 2 positive real-valued parameters, i.e., a vector α = (α₁,..., α_K) ∈ ℝ^K₊.
- **\blacksquare** The probability density function is defined on the K simplex

$$\Delta_{\mathcal{K}} = \left\{ \boldsymbol{\theta} = (\theta_1, \dots, \theta_{\mathcal{K}})^\top \, | \, \theta_1, \dots, \theta_{\mathcal{K}} \geq 0, \, \sum_{k=1}^{\mathcal{K}} \theta_k = 1 \right\}$$

and given as follows:

$$p(\boldsymbol{ heta} \mid \boldsymbol{lpha}) = p(heta_1, \dots, heta_K \mid \boldsymbol{lpha}) = rac{1}{\mathbb{B}(\boldsymbol{lpha})} \prod_{k=1}^K heta_k^{lpha_k - 1},$$

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where the normalisation constant is the multivariate Beta function.

In Bayesian statistics, the Dirichlet distribution is commonly used as the conjugate prior of the multinomial distribution.

Dirichlet distribution with parameters $\alpha = (\alpha_1, \alpha_2) = (1, 1)$, (5, 5), (3, 6), (3, 12).



Predicting a Dirichlet distribution



Several authors have proposed the minimisation of an empirical loss of the form

$$L = \frac{1}{N} \sum_{n=1}^{N} \ell_2 (Q^{(n)}, y^{(n)}) = \frac{1}{N} \sum_{n=1}^{N} \mathbb{E}_{\theta \sim Q} \ell_1 \left(\theta, y^{(n)}\right) ,$$

where $Q^{(n)}$ is the level-2 prediction for the instance $\mathbf{x}^{(n)}$.

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- Examples of level-1 loss include cross entropy (Charpentier *et al.*, 2020) and Brier score (Sensoy *et al.*, 2018).
- Besides, a regularised version has been proposed:

$$L = \frac{1}{N} \sum_{n=1}^{N} \underbrace{\mathbb{E}_{\theta \sim Q} \ell_{1}(\theta, y) + \lambda d_{KL}\left(Q^{(n)}, Q_{0}\right)}_{\ell_{2}\left(Q^{(n)}, y^{(n)}\right)}$$

A negative result

Informally, we define a level-2 loss function ℓ_2 as **appropriate** if the following holds for the empirical loss minimiser

$$Q^{(N)} = rgmin_Q rac{1}{N} \sum_{n=1}^N \ell_2\left(Q, y^{(n)}
ight)$$

on any i.i.d. observational data sequence $y^{(1)}, y^{(2)}, \ldots$ with $y^{(i)} \sim \boldsymbol{\theta}^*$:

- (A1) The learner's uncertainty gradually decreases (in expectation) with increasing sample size N, in terms of a suitable uncertainty measure U.
- (A2) In the limit $N \to \infty$, all epistemic uncertainty disappears and $Q^{(N)} \to \delta_{\theta^*}$.

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 (A2) In the limit N → ∞, all epistemic uncertainty disappears and Q^(N) → δ_{θ*}.
- Bengs *et al.* (2022) formally prove that a loss minimisation approach using a level-2 loss as specified above does not lead to an appropriate level-2 loss.

Generalised uncertainty calculi



Credal uncertainty representation



ENSEMBLE-BASED UNCERTAINTY QUANTIFICATION: BAYESIAN VERSUS CREDAL INFERENCE

A PREPRINT

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December 13, 2021

Possibilistic uncertainty representation
Possibilistic uncertainty representation

Senge *et al.* (2014) formalise the following principle in a mathematical way: Given a query \mathbf{x}_q , an outcome $y \in \mathcal{Y}$ is a **plausible** candidate if there exists a **plausible** hypothesis $h \in \mathcal{H}$ (compatible with the data) such that y is **strongly supported** by h (in the sense that $p(y | \mathbf{x}_q, h)$ is high).

Possibilistic uncertainty representation

- Senge et al. (2014) formalise the following principle in a mathematical way: Given a query x_q, an outcome y ∈ 𝔅 is a plausible candidate if there exists a plausible hypothesis h ∈ 𝕂 (compatible with the data) such that y is strongly supported by h (in the sense that p(y | x_q, h) is high).
- Thus, they induce a graded (fuzzy) set of plausible candidate hypotheses instead of a single one, and consider the plausible outcomes under these candidates.



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- Related to this is the "closed world" assumption, which is often violated in practice, e.g., in the case of OOD data.
- Usefulness of generalized uncertainty calculi?

More on this topic ...

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Aleatoric and epistemic uncertainty in machine learning: an introduction to concepts and methods

Eyke Hüllermeier 🖂 & Willem Waegeman

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Abstract

The notion of uncertainty is of major importance in machine learning and constitutes a key element of machine learning methodology. In line with the statistical tradition, uncertainty has long been perceived as almost synonymous with standard probability and probabilistic predictions. Yet, due to the steadily increasing relevance of machine learning for practical applications and related issues such as safety requirements, new problems and challenges have recently been identified by machine learning scholars, and these problems may call for new methodological developments. In particular, this includes the importance of distinguishing between (at least) two different types of uncertainty, often referred to as *aleatoric* and *epistemic*. In this paper, we provide an introduction to the topic of uncertainty in machine learning as well as an overview of attempts so far at handling uncertainty in general and formalizing this distinction in particular.

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Epistemic uncertainty sampling

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How to measure uncertainty in uncertainty sampling for active learning

Vu-Linh Nguyen, Mohammad Hossein Shaker & Eyke Hüllermeier 🖂

Machine Learning (2021) | Cite this article 1033 Accesses | 1 Altmetric | Metrics

Abstract

Various strategies for active learning have been proposed in the machine learning literature. In uncertainty sampling, which is among the most popular approaches, the active learner sequentially queres the label of those instances for which its current prediction is maximally uncertain. The predictions as well as the measures used to quantify the degree of uncertainty, such as entropy, are traditionally of a probabilistic nature. Yet, alternative approaches to capturing uncertainty in machine learning, alongide with corresponding uncertainty measures, have been proposed in recent years. In particular, some of these measures seek to distinguish different sources and to separate different types of uncertainty as has the reducible (epistemic) and the irreducible (aleatoric) part of the total uncertainty in a prediction. The goal of this paper is to laborate on the usefulness of such measures for uncertainty sampling, and to compare their performance in active learning. To this end, we instantiate uncertainty sampling with different measures, analyze the properties of the sampling strategies thus obtained, and compare them in an experimental study.



Sampling in epistemically uncertain regions of the instance space is potentially more useful than sampling in aleatorically uncertain regions ...

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- Suppose the distribution is concentrated in a "normal" range, so that \mathcal{T} takes values in $A = [t_l, t_u] \subset \mathbb{R}$ with high probability 1δ , where δ is the significance level.

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- Suppose the distribution is concentrated in a "normal" range, so that T takes values in $A = [t_l, t_u] \subset \mathbb{R}$ with high probability 1δ , where δ is the significance level.
- **That is, assuming** H_0 , it is unlikely to observe values $T \notin A$.
- If this nevertheless happens, i.e., $T \in R = \mathbb{R} \setminus A$, then H_0 is rejected and H_1 accepted.

Evaluation: accuracy-rejection curves

Evaluation: accuracy-rejection curves

Reject test instances for which (total, aleatoric, epistemic) uncertainty exceeds a certain threshold, measure accuracy on the remaining ones.



Facets of uncertainty

Facets of uncertainty

Randomness, imprecision, inconsistency, ambiguity, vagueness, fuzziness, ...



"Fuzziness is orthogonal to probability theory – it focuses on the ambiguity of describing events, rather than the uncertainty about the occurrence or non-occurrence of events." J

Judea Pearl (2000)

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- $\blacksquare H_0: \theta = \frac{1}{2}, H_1: \theta \neq \frac{1}{2}$
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- $\mathcal{D} = \{x_1, \ldots, x_N\}$, outcomes of N coin flips
- **T** = number of heads in \mathcal{D} .
- **Distribution under** H_0 is binomial:

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Take $R = \{0, 1, \dots, r\} \cup \{N - r, \dots, N\}$, with r the largest number such that

$$P(T \in R) = \sum_{k=0}^{r} B(k, 1/2) + \sum_{k=N-r}^{N} B(k, 1/2) \le \delta.$$





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Excursus: Hypothesis testing

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- **Type-II error:** not rejecting *H*₀ although it is false—the probability of this error depends on the (then unknown) ground truth.
- Also note that **nothing** can be concluded from the test in case H_0 is not rejected.
- In particular, not rejecting H_0 does not mean one can "accept" it:

$$P\left(\theta=\frac{1}{2}\mid T\in\{6,\ldots,14\}\right)=??$$

■ A (finite) sequence of random variables $X_1, X_2, ..., X_N$ is exchangeable, if the following holds for any permutation $\sigma : [N] \longrightarrow [N]$: the joint probability distribution of the random variables is the same as the joint distribution of the permuted sequence $X_{\sigma(1)}, X_{\sigma(2)}, ..., X_{\sigma(N)}$.

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Exchangeability is weaker than the i.i.d. assumption (the latter implies the former but not the other way around).

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- Inductive conformal prediction (ICP) is an alternative that is computationally less expensive.
- In the split-conformal prediction variant, ICP splits the training data \mathcal{D} into
 - proper training data \mathcal{D}_T of size N M,
 - calibration data $\mathcal{D}_C = \{(\mathbf{x}_j, y_j)\}_{j=1}^M$ of size M < N.

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Assuming \mathcal{D} to be an i.i.d. sample, and (\mathbf{x}_q, y_q) sampled from the same distribution, the following estimate is provably correct:

$$1-\epsilon \leq P(y_q \in Y(\boldsymbol{x}_q)) \leq 1-\epsilon + \frac{1}{M+1}$$