## Dropout as a Bayesian Approximation

appliedAI Seminar — Uncertainty Quantification in Deep Learning

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## INITIATIVE FOR APPLIED ARTIFICIAL INTELLIGENCE

1. Motivation

2. Quick intro to Uncertainty Quantification

3. Recap of Dropout in Neural Networks

4. MC Dropout for Uncertainty Quantification in Deep Learning

# Motivation

#### Why do we want to quantify the uncertainty in our models?



Figures showing the true value (red) and the predictive mean (blue) including  $\pm 2\sigma$  for a model evaluated on the Mauna Loa CO<sub>2</sub> concentration dataset [4]. The blue shades indicate half a std. deviation each.

**Uncertainty Quantification** 

#### Uncertainty is devided into <u>aleatoric</u> and epistemic uncrtainty

#### Aleatoric uncertainty

- Stochastic uncertainty or unknowns, responsible for different outcomes of the same experiment
- Often due to insufficient measurement quality
- Does not preclude the existence of the "unknowns"
- Uncertainty trapped "in the data"
- Methods: Quantile regression, mixture of densities, max. likelihood est.

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### Epistemic uncertainty

- Systematic uncertainty, due to possibly available information but practically not used
- Inaccuracies in measurements, unused data / features, neglecting effects
- Captured by the model due to design decisions, data used, etc.
- Methods: Bayesian Neural Networks, MC dropout or deep ensembles

#### Illustration of aleatoric uncertainty



Based on the current information (years in the job), the increasing variance in ML experience cannot be explained.

## Illustration of epistemic uncertainty



While the model returns point estimates, it is uncertain about its predictions, especially further in the future.

**Recap of Dropout** 

### Dropping our neurons to prevent overfitting



Illustration of a standard neural network vs. one with dropping out nodes [6].

Where the keep probability is p and layers are denoted by  $l \in \{0, ..., L-1\}$  [6].

- Reduce risk of overfitting
- Model ensemble without optimizing hundreds of models
- Activate all parts of the model

### Dropout is primarly used to prevent overfitting

- Reduce risk of overfitting
- Model ensemble without optimizing hundreds of models
- Activate all parts of the model



Dropout improves performance on the test data for different architectures and configurations [6].

### Dropout is switched offduring inference — all nodes are active



As dropping nodes is turned off during inference, the weights need to be scaled to not get larger than expected [6].

Monte Carlo Dropout

#### Q: How do we get the model's uncertainty?



Figures showing the true value (red) and the predictive mean (blue) including  $\pm 2\sigma$  for a model evaluated on the Mauna Loa CO<sub>2</sub> concentration dataset [4]. The blue shades indicate half a std. deviation each.

### Retrieving the predictive mean and variance

Given a model  $\mathcal{M}$  which was trained with dropout and a *keep probability* of p. Generate  $\{(\hat{y}_i, \mathbf{x})\}_{i=1}^N$  by **keeping dropout on**.

Estimators for  $\mathbb{E}(y)$ , Var(y) read

$$\mathbb{E}(\mathbf{y}) \approx \frac{1}{N} \sum_{i=1}^{N} \hat{y}_i(\mathbf{x})$$

$$\operatorname{Var}(\mathbf{y}) \approx \tau^{-1} \mathbf{I} + \frac{1}{N} \sum_{i=1}^{N} \hat{y}_i(\mathbf{x}) \hat{y}_i(\mathbf{x})^T - \mathbb{E}(\mathbf{y}) \mathbb{E}(\mathbf{y})^T$$
(3)

Where  $\tau = \frac{l^2 p}{2N\lambda}$  the GP precision;  $\tau, p$  obtained via hyperparameter search [5].

**Algorithm 1:** Implementation of Monte Carlo Dropout by using dropout during inference as well and computing statistics of the samples from the predictive distribution.

```
1: \mathcal{M} \leftarrow \mathcal{M}.train()

2: \hat{y}_i \leftarrow \mathcal{M}(\mathbf{x}) \forall i \in [1, ..., N]

3:

4: \hat{\mathbb{E}}(\mathbf{y}) \leftarrow \hat{\mathbf{y}}.mean()

5: \hat{\sigma}(\mathbf{y}) \leftarrow \hat{\mathbf{y}}.std()

6:

7: return \hat{\mathbb{E}}(\mathbf{y}), \hat{\sigma}(\mathbf{y})
```

**•** "Averaging over forward passes is equivalent to Monte Carlo integration over a Gaussian Process posterior approximation."

 $\rightarrow$  We'll familiarize with the idea **but** I refer to [3, 4] for a detailed discussion.

**•** "Averaging over forward passes is equivalent to Monte Carlo integration over a Gaussian Process posterior approximation."

→ We'll familiarize with the idea **but** I refer to [3, 4] for a detailed discussion. **Target:** predictive distribution for a new observation  $x^*$ 

$$p(y^*|\mathbf{x}^*, \mathbf{X}, \mathbf{Y}) = \int p(y^*|\mathbf{x}^*, \omega) p(\omega|\mathbf{X}, \mathbf{Y}) d\omega$$

$$\approx \int p(y^*|\mathbf{x}^*, \omega) q_{\theta^*}(\omega) d\omega, \quad \theta^* = \underset{\theta}{\operatorname{argmin}} \operatorname{KL}(q_{\theta}(\omega)|p(\omega|\mathbf{X}, \mathbf{Y}))$$
(4)

## Finding $q_{\theta^*} \Rightarrow$ Minimizing the KL-Divergence

 $\P$  Minimizing KL-Div.  $\Leftrightarrow$  maximizing log evidence lower bound [1].

$$\begin{split} \log p(\mathbf{Y}|\mathbf{X}) &= \log \int p(\mathbf{Y}|\mathbf{X},\omega)p(\omega)d\omega \\ &= \log \int p(\mathbf{Y}|\mathbf{X},\omega)p(\omega)\frac{q_{\theta}(\omega)}{q_{\theta}(\omega)}d\omega \\ &= \log \left(\mathbb{E}_{q_{\theta}}\frac{p(\mathbf{Y}|\mathbf{X},\omega)p(\omega)}{q_{\theta}(\omega)}\right) \end{split}$$
(5)  

$$\begin{split} \text{Jensen's ineq.} \\ &\geq \\ \mathbb{E}_{q_{\theta}}\log \left(\frac{p(\mathbf{Y}|\mathbf{X},\omega)p(\omega)}{q_{\theta}(\omega)}\right) \\ L_{\text{VI}} &= \int q_{\theta}(\omega)\log(p(\mathbf{Y}|\mathbf{X},\omega))d\omega - \text{KL}(q_{\theta}|p(\omega)) \end{split}$$

Using MC integration to approximate the integral with one sample  $\omega \sim q_{\theta}$  and optimizing w.r.t.  $\theta$ :

$$\hat{L}(\theta) = \log p(\mathbf{Y}|\mathbf{X}, \hat{\omega}) - \mathsf{KL}(q_{\theta}|p(\omega))$$
(6)

 $\Rightarrow q_{\theta}(\omega)$  is a distribution over weights of a network. Thus, define  $q_{\theta}(\omega)$  s.t. weights can be turned off.

$$q_{\theta^*} = \prod_i q_{\mathsf{M}_i}(\mathsf{W}_i), \quad i \in [1, \dots, L]$$

$$q_{\mathsf{M}_i}(\mathsf{W}_i) = \mathsf{M}_i \operatorname{diag}(z_{i,j}) \qquad (7)$$

$$z_{i,j} \sim \mathcal{B}(p_i)$$

$$\mathsf{W}_i \sim q_{\mathsf{M}_i}(\mathsf{W}_i)$$

- Requires large number of samples making inference slow (>  $10^3$  in [4])
- Inference time is crucial in a lot of applications

- "Succession" of the original Monte Carlo Dropout
- Preserves advantages of Bayesian NN without being slower than non-Bayesian NN [2]
- Idea: approximate the expected value and variance of the MC Dropout per layer

### Approximation via Moment Propagation



Illustration of Moment Propagation through dropout, fully-connected and ReLU layers [2]. The expectation and variance *flow* through the network during a single forward pass

Dropout Layer per layer with nodes *i* 

$$E^{i} = E^{i}_{in} \cdot p$$

$$V^{i} \stackrel{X \perp P}{=} V^{i}_{in} \cdot p(1-p) + V^{i}_{in}p^{2} + (E^{i}_{in})^{2} \cdot p(1-p)$$
(8)

**Dropout Layer** per layer with nodes *i* 

Dense Layer

$$E' = E'_{in} \cdot p$$

$$V^{i} \stackrel{X \perp Y}{=} V^{i}_{in} \cdot p(1-p) + V^{i}_{in}p^{2} + (E^{i}_{in})^{2} \cdot p(1-p)$$
(8)

$$E^{i} = \sum_{j}^{N_{\text{in}}} w_{ji} E^{j}_{\text{in}} + b^{i}$$

$$V^{i} = \sum_{j}^{N_{\text{in}}} w^{2}_{ji} V^{j}_{\text{in}}$$
(9)

#### Comparison of Single Shot MC and MC Dropout



Visual comparison of the two methods using MC Dropout. The models were trained on data in the range of [-3, 19]. In the upper figure, the predictions are shown. In the lower panel, the uncertainty is included, showing similar results [2].

### Comparison of Single Shot MC and MC Dropout ctd.

DATASET	Ν	Q	TEST RMSE		TEST NLL		TEST RT [S]	
			MC	MP	MC	MP	MC	MP
BOSTON	506	13	$3.14 \pm 0.20$	$3.10 \pm 0.20$	$2.57 \pm 0.07$	$2.56 \pm 0.08$	$2.51 \pm 0.03$	$0.04 \pm 0.00$
Concrete	1,030	8	$5.46 \pm 0.12$	$5.40 \pm 0.12$	$3.12 \pm 0.02$	$3.13 \pm 0.03$	$3.37 \pm 0.04$	$0.04 \pm 0.00$
Energy	768	8	$1.65 \pm 0.05$	$1.61 \pm 0.05$	$1.95 \pm 0.04$	$2.01 \pm 0.04$	$2.84 \pm 0.03$	$0.04 \pm 0.00$
Kin8nm	8,192	8	$0.08 \pm 0.00$	$0.08 \pm 0.00$	$-1.10 \pm 0.01$	$-1.11 \pm 0.01$	$7.37 \pm 0.06$	$0.04 \pm 0.00$
NAVAL	11,934	16	$0.00\pm\!0.00$	$0.00\pm\!0.00$	$-4.36 \pm 0.01$	$-3.60 \pm 0.01$	$9.69 \pm 0.11$	$0.04 \pm 0.00$
POWER	9,568	4	$4.05 \pm 0.04$	$4.04 \pm 0.04$	$2.82 \pm 0.01$	$2.84 \pm 0.01$	$6.85 \pm 0.07$	$0.04 \pm 0.00$
Protein	45,730	9	$4.42 \pm 0.03$	$4.41 \pm 0.02$	$2.90 \pm 0.00$	$2.91 \pm 0.00$	$31.38 \pm 0.09$	$0.05 \pm 0.00$
WINE	1,599	11	$0.63 \pm 0.01$	$0.63 \pm 0.01$	$0.95 \pm 0.01$	$0.95 \pm 0.01$	$4.78 \pm 0.01$	$0.04 \pm 0.00$
YACHT	308	6	$2.93 \pm 0.22$	$2.91 \pm 0.26$	$2.35 \pm 0.07$	$2.11 \pm 0.07$	$2.01 \pm 0.01$	$0.04 \pm 0.00$

Benchmarking both approaches on the UCI benchmarking datasets where  $N = 10^4$  forward passes were used for the MC Dropout model. Both methods show similar results on the test data while the approach of Moment Propagation has a significant advantage w.r.t. inference time [2].

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