

# Computer-intensive Estimation and Prediction for Time Series and Deep Neural Networks

Defense Talk

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1. Prediction inference of Non-linear Autoregressive models
2. Deep Model-free generative prediction method for regression
3. Scalable subsampling for DNN training

# Prediction inference of Non-linear Autoregressive models<sup>1</sup>

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<sup>1</sup>This part is based on:

- Wu, K. and Politis, D.N., Bootstrap Prediction Inference of Nonlinear Autoregressive Models, *Journal of Time Series Analysis* 2024, 45, 800-822.
- Wu, K., Non-parametric Forward Bootstrap on Predicting Non-linear Time Series: Consistency, Pertinence and Debiasing, *Stats* 2023, 6(3), 839-867.



- **Time series** is a (discrete-time) stochastic process, i.e.,  $\{X_t, t \in \mathbb{Z}\}$ . Its realization is called time series data, e.g., heights of ocean tides, and counts of sunspots.

- **Time series** is a (discrete-time) stochastic process, i.e.,  $\{X_t, t \in \mathbb{Z}\}$ . Its realization is called time series data, e.g., heights of ocean tides, and counts of sunspots.
- **Prediction inference** is about determining Optimal Predictor (OP), usually in  $L_2$  or  $L_1$  sense, and Prediction Interval (PI), percentile or centered version, of future value  $X_{T+k}$ ,  $k \geq 1$ , based on observed  $\{X_0, \dots, X_T\}$ . We are concerned about the Coverage Rate (CVR) and Length (LEN) of PI.

# Two situations

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### Simple case:

If  $\{X_0, \dots, X_T\}$  are *i.i.d.*. Take sample mean and sample median to be  $L_2$  and  $L_1$  OPs, respectively. Rely on sample quantile values to build PIs.



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### Time series model:

We assume that the time series data is generated by some underlying mechanism:

$$X_t = G(X_{t-p}, \epsilon_t),$$

where:

- $G(\cdot, \cdot)$  could be any suitable linear/non-linear function that makes the time series have desired property.
- $\epsilon_t \sim F_\epsilon$  is called innovation and assumed to be *i.i.d.* with appropriate moments and independent with  $X_{t-i}$ ,  $i \geq 1$ .
- $X_{t-p}$  represents  $\{X_{t-1}, \dots, X_{t-p}\}$ .

# Monte Carlo (MC) simulation for multi-step ahead prediction

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Apply Monte Carlo (MC) simulation to do predictions:

- 1 Simulate  $\{\epsilon_{T+1}^{(i)}, \dots, \epsilon_{T+k}^{(i)}\}_{i=1}^M$  from  $F_\epsilon$ .
- 2 Compute pseudo  $\{X_{T+k}^{(i)}\}_{i=1}^M$ , i.e.,  $X_{T+j}^{(i)} = G(X_{T+j-p}, \epsilon_{T+j}^{(i)})$ , for  $j = 1, \dots, k$ .
- 3 Take sample mean and median of  $\{X_{T+k}^{(i)}\}_{i=1}^M$  to approximate optimal predictors, respectively. Take corresponding quantile values to approximate PIs with arbitrary coverage rates. We call such type of PI Simulation PI (SPI).

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**Limitation:** In practice, model information is generally not known to participators. Thus, this prediction is *Oracle*.

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Apply Bootstrap to do predictions:

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- 2 Compute pseudo  $\{\widehat{X}_{T+k}^{(i)}\}_{i=1}^M$  iteratively, i.e.,  $\widehat{X}_{T+j}^{(i)} = \widehat{G}(X_{T+j-p}, \hat{\epsilon}_{T+j}^{(i)})$ , for  $j = 1, \dots, k$ .
- 3 Take sample mean and median of  $\{\widehat{X}_{T+k}^{(i)}\}_{i=1}^M$  to approximate optimal predictors, respectively. Take corresponding quantile values to approximate PIs with arbitrary coverage rates. We call such type of PI Quantile PI (QPI).

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**Limitation:** In practice with finite samples, this Bootstrap-based PI suffers undercoverage.

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- 3 Re-estimate model to get  $\widehat{G}^*(\cdot, \cdot)$  with  $\{X_0^*, \dots, X_T^*\}$ ; Re-define  $\{X_{T-p+1}^* = X_{T-p+1}, \dots, X_T^* = X_T\}$ . Then do the bootstrap prediction with  $\widehat{G}^*(\cdot, \cdot)$  and  $\widehat{F}_\epsilon$  to get  $\widehat{X}_{T+k}^*$ . Record the predictive root  $X_{T+k}^* - \widehat{X}_{T+k}^*$  in the bootstrap world.

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- 4 Repeat the above process  $M$  times, collect  $M$  predictive roots and take its empirical distribution to approximate the distribution of  $X_{T+k} - \widehat{X}_{T+k}$ .
- 5 The  $(1 - \alpha)100\%$  PI for  $X_{T+k}$  centered at  $\widehat{X}_{T+k}$  can be approximated by  $[\widehat{X}_{T+k} + q(\alpha/2), \widehat{X}_{T+k} + q(1 - \alpha/2)]$ , where  $q(\alpha)$  is the  $\alpha$ -quantile of the empirical distribution of  $X_{T+k}^* - \widehat{X}_{T+k}^*$ .

# Pertinent Prediction Interval (PPI)

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$$2 \quad \sup_a |\mathbb{P}(\tau_T A_m^* \leq a) - \mathbb{P}(\tau_T A_m \leq a)| \xrightarrow{P} 0,$$

For example, we assume that we can decompose  $G(X_{t-p}, \epsilon_t)$  as  $M(X_{t-p}) + \epsilon_t$ ;  
 $A_m^* = \widehat{M}^*(\mathbf{x}) - \widehat{M}(\mathbf{x})$ ;  $A_m = \widehat{M}(\mathbf{x}) - M(\mathbf{x})$ .

# Parametric Non-linear Autoregressive (NLAR) models

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First, we consider the case that we can decompose  $G(X_{t-p}, \epsilon_t)$  as a parametric non-linear model<sup>2</sup>:

$$X_t = G(X_{t-1}, \epsilon_t) = m(X_{t-1}, \theta_1) + \sigma(X_{t-1}, \theta_2)\epsilon_t,$$

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where:

- $m(\cdot)$  is the mean function which is Lipschitz continuous w.r.t. the first and second arguments for their domain, respectively.
- $\sigma(\cdot)$  is the positive and bounded variance function which is Lipschitz continuous w.r.t. the first and second arguments for their domains, respectively.
- $\theta_1 \in \Theta_1$  and  $\theta_2 \in \Theta_2$ , where  $\Theta_1$  and  $\Theta_2$  are all bounded sets in  $\mathbb{R}^d$ .
- For  $\epsilon_t$ , it is mean zero and variance 1;  $f_\epsilon(\cdot)$  is continuous and everywhere positive.

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## Two-step estimation process

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$$\widehat{\theta}_1 = \arg \min_{\vartheta \in \Theta_1} L_T(\vartheta) = \arg \min_{\vartheta \in \Theta_1} \frac{1}{T} \sum_{t=1}^T (X_t - m(X_{t-1}, \vartheta))^2$$

## Two-step estimation process

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•

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•

$$\widehat{\theta}_2 = \arg \min_{\vartheta \in \Theta_2} K_T(\vartheta, \widehat{\theta}_1) = \arg \min_{\vartheta \in \Theta_2} \left| \frac{1}{T} \sum_{t=1}^T \left( \frac{X_t - m(X_{t-1}, \widehat{\theta}_1)}{\sigma(X_{t-1}, \vartheta)} \right)^2 - 1 \right|.$$





# Consistency of OP and asymptotic validity of QPI

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## Theorem 1: Consistency of prediction

For  $k \geq 1$  we have:

$$\sup_{|x| \leq c_T} |F_{X_{T+k}^* | X_T, \dots, X_0}(x) - F_{X_{T+k} | X_T}(x)| \xrightarrow{p} 0,$$

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where

- $X_{T+k}^* = \mathcal{G}(X_T; \hat{\epsilon}_{T+1}^*, \dots, \hat{\epsilon}_{T+k}^*; \widehat{\theta})$ . This is computed by  $X_{T+i}^* = m(X_{T+i-1}^*, \widehat{\theta}_1) + \sigma(X_{T+i-1}^*, \widehat{\theta}_2) \hat{\epsilon}_{T+i}^*$  iteratively for  $i = 1, \dots, k$ . Similar for  $X_{T+k}$ .
- $\{\hat{\epsilon}_i^*\}_{i=T+1}^{T+k}$  are *i.i.d.*  $\sim \widehat{F}_\epsilon$ .
- $c_T$  is an appropriate sequence converges to infinity as  $T$  converges to infinity.
- $F_{X_{T+k}^*|X_T, \dots, X_0}(x)$  is the distribution of  $k$ -step ahead future value in the bootstrap world, i.e., conditional on all observed data.
- $F_{X_{T+k}|X_T}(x)$  is the distribution of  $k$ -step ahead future value in the real world.

# Estimation inference of $\widehat{\theta}_1$ and $\widehat{\theta}_2$ , $\widehat{\theta}_1^*$ and $\widehat{\theta}_2^*$

## Theorem 2: Estimation inference

Based on the realization  $\{X_0, \dots, X_T\} \in \Omega_T$ , where  $\mathbb{P}((X_0, \dots, X_T) \notin \Omega_T) = o(1)$  as  $T \rightarrow \infty$ , under other suitable assumptions, we have:

$$\begin{aligned}\sqrt{T}(\widehat{\theta}_1 - \theta_1) &\xrightarrow{d} N(0, B_1^{-1} \Omega_1 B_1^{-1}); \quad \sqrt{T}(\widehat{\theta}_2 - \theta_2) \xrightarrow{d} N(0, B_2^{-1} \Omega_2 B_2^{-1}); \\ \sqrt{T}(\widehat{\theta}_1^* - \widehat{\theta}_1) &\xrightarrow{d} N(0, B_1^{-1} \Omega_1 B_1^{-1}); \quad \sqrt{T}(\widehat{\theta}_2^* - \widehat{\theta}_2) \xrightarrow{d} N(0, B_2^{-1} \Omega_2 B_2^{-1});\end{aligned}$$

where

- $\Omega_1 = 4 \cdot \mathbb{E}(\sigma(X_0, \theta_2) R_1 \sigma(X_0, \theta_2)); B_1 = 2 \cdot \mathbb{E}(\nabla \phi(X_0, \theta_1) (\nabla \phi(X_0, \theta_1))^\top); R_1 = \nabla \phi(X_0, \theta_1) \nabla \phi(X_0, \theta_1)^\top$ ; here  $\nabla$  is the gradient operator w.r.t.  $\theta_1$ .
- $\Omega_2 = 4 \cdot \mathbb{E}(B_3 R_2 B_3^\top); B_3 = \mathbb{E}(\nabla g(X_1, X_0, \theta_2, \theta_1)); R_2 = (g(X_1, X_0, \theta_2, \theta_1) - 1)^2$ ;  
 $B_2 = 2 \cdot (\mathbb{E}(\nabla g(X_1, X_0, \theta_2, \theta_1)) \cdot (\mathbb{E}(\nabla g(X_1, X_0, \theta_2, \theta_1))^\top); g(X_1, X_0, \theta_2, \theta_1) = \left( \frac{X_1 - \phi(X_0, \theta_1)}{\sigma(X_0, \theta_2)} \right)^2$ ; here  $\nabla$  is the gradient operator w.r.t.  $\theta_2$ .

# Non-parametric NLAR models

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When the parametric format is unknown, we assume that we only know the data-generating mechanism of time series consists of two parts:

$$X_t = G(X_{t-1}, \epsilon_t) = m(X_{t-1}) + \sigma(X_{t-1})\epsilon_t.$$

$$\widetilde{m}_h(x) = \frac{\sum_{t=1}^T K(\frac{x-X_{t-1}}{h})X_t}{\sum_{t=1}^T K(\frac{x-X_{t-1}}{h})} \quad \text{and} \quad \widetilde{\sigma}_h(x) = \frac{\sum_{t=1}^T K(\frac{x-X_{t-1}}{h})(X_t - \widetilde{m}_h(X_{t-1}))^2}{\sum_{t=1}^T K(\frac{x-X_{t-1}}{h})};$$

## Theorem 3: QPI and PPI of non-parametric prediction

Let  $\widehat{m}_g(x)$  and  $\widehat{\sigma}_g(x)$  be estimated mean and variance functions to generate bootstrap series in the bootstrap world. With the under-smoothing debiasing strategy, i.e., we take  $g = h$  and take a bandwidth rate satisfying  $hT^{1/5} \rightarrow 0$ . The QPI and PPI are still possible with the main prediction algorithm.

# Simulation for parametric prediction approach

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**Simulation model** (Threshold model):

$$X_t = (0.5 \cdot X_{t-1} + 0.2 \cdot X_{t-2} + 0.1 \cdot X_{t-3})I(X_{t-1} \leq 0) + (0.8 \cdot X_{t-1})I(X_{t-1} > 0) + \epsilon_t; \epsilon_t \sim N(0, 1).$$

**Simulation setting:**

We take the number of bootstrap times  $M = 1000$ . We repeat simulations  $N = 5000$  times. We take  $\alpha = 0.05$ .

**Simulation measurement:**

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$$\text{CVR of the } k\text{-th step ahead prediction} = \frac{1}{N} \sum_{n=1}^N I_{X_{n,k} \in [L_{n,k}, U_{n,k}]}, \text{ for } k = 1, \dots, 5.$$

•

$$\text{LEN of the } k\text{-th step ahead PI} = \frac{1}{N} \sum_{n=1}^N (U_{n,k} - L_{n,k}), \text{ for } k = 1, \dots, 5,$$

where  $[L_{n,k}, U_{n,k}]$  and  $X_{n,k}$  represent  $k$ -th step ahead prediction intervals and the true future value in the  $n$ -th replication, respectively.



# Simulation results

Threshold Model:		$X_t = (0.5 \cdot X_{t-1} + 0.2 \cdot X_{t-2} + 0.1 \cdot X_{t-3})I(X_{t-1} \leq 0) + (0.8 \cdot X_{t-1})I(X_{t-1} > 0) + \epsilon_t, \epsilon_t \sim N(0, 1)$								
$T = 400$	CVR for each step					LEN for each step				
	1	2	3	4	5	1	2	3	4	5
QPI-f	0.9420	0.9506	0.9468	0.9444	0.9372	3.88	4.68	5.11	5.40	5.58
QPI-p	0.9462	0.9512	0.9502	0.9474	0.9428	3.92	4.72	5.16	5.45	5.64
$L_2$ -PPI-f	0.9446	0.9510	0.9486	0.9470	0.9408	3.90	4.71	5.15	5.44	5.63
$L_2$ -PPI-p	0.9466	0.9542	0.9516	0.9494	0.9434	3.94	4.75	5.20	5.49	5.69
$L_1$ -PPI-f	0.9448	0.9518	0.9478	0.9468	0.9402	3.90	4.71	5.15	5.44	5.62
$L_1$ -PPI-p	0.9470	0.9544	0.9500	0.9486	0.9436	3.94	4.75	5.20	5.49	5.68
SPI	0.9446	0.9534	0.9508	0.9510	0.9454	3.90	4.71	5.16	5.46	5.65
$T = 100$										
QPI-f	0.9270	0.9304	0.9294	0.9272	0.9250	3.81	4.57	4.98	5.23	5.40
QPI-p	0.9370	0.9412	0.9368	0.9372	0.9372	3.98	4.76	5.19	5.46	5.63
$L_2$ -PPI-f	0.9358	0.9352	0.9338	0.9314	0.9298	3.95	4.71	5.13	5.40	5.59
$L_2$ -PPI-p	0.9454	0.9454	0.9444	0.9430	0.9418	4.10	4.90	5.34	5.63	5.83
$L_1$ -PPI-f	0.9364	0.9360	0.9336	0.9310	0.9304	3.95	4.71	5.13	5.39	5.58
$L_1$ -PPI-p	0.9450	0.9456	0.9432	0.9422	0.9412	4.11	4.90	5.33	5.62	5.81
SPI	0.9446	0.9472	0.9498	0.9474	0.9478	3.90	4.71	5.16	5.46	5.65
$T = 50$										
QPI-f	0.8980	0.9054	0.9018	0.8950	0.8926	3.66	4.47	4.87	5.14	5.38
QPI-p	0.9260	0.9314	0.9272	0.9218	0.9212	4.05	4.97	5.42	5.74	5.99
$L_2$ -PPI-f	0.9340	0.9268	0.9214	0.9164	0.9152	4.22	5.10	5.86	6.89	8.97
$L_2$ -PPI-p	0.9522	0.9478	0.9404	0.9400	0.9376	4.60	5.57	6.36	7.33	9.03
$L_1$ -PPI-f	0.9338	0.9268	0.9194	0.9144	0.9130	4.23	5.09	5.82	6.79	8.71
$L_1$ -PPI-p	0.9522	0.9482	0.9384	0.9378	0.9356	4.61	5.55	6.30	7.20	8.71
SPI	0.9494	0.9448	0.9464	0.9458	0.9462	3.90	4.71	5.16	5.46	5.65

Note: “-f” and “-p” represent fitted and predictive residuals, respectively. “ $L_2$ ” and “ $L_1$ ” represent the center of PPI is  $L_2$  and  $L_1$  OP, respectively.

## Deep Model-free generative prediction method for regression<sup>3</sup>

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<sup>3</sup>This part is based on:

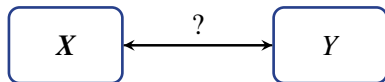
- Wu, K. and Politis, D.N., Deep Limit Model-free Prediction in Regression. (*Submitted to ACM/IMS Journal of Data Science*)



# Regression analysis

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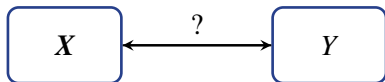
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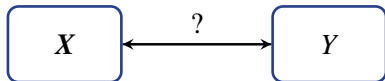
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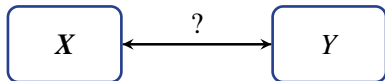
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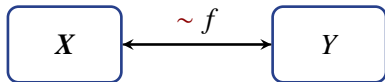


## Model as bridge

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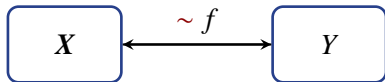
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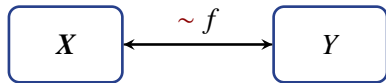
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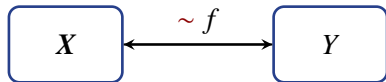
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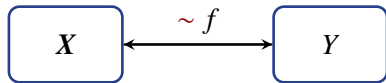
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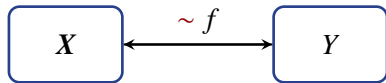
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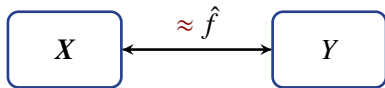
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In practice, we estimate  $f(\cdot)$  by  $\hat{f}(\cdot)$  based on sample  $\{X_i, Y_i\}_{i=1}^n$ :<sup>5</sup>



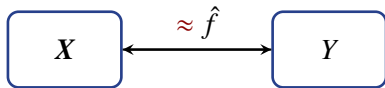
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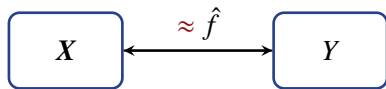
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Therefore, we need to quantify the estimation accuracy, e.g., by Confidence Interval (CI).

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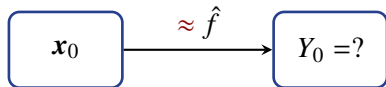
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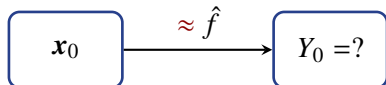
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To quantify the prediction accuracy, we build Prediction Interval (PI). However, it usually requires the normality assumption or it suffers the undercoverage in the finite sample cases.

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- Even a complicated  $Y = f(X) + g(X) \cdot \varepsilon$ , where  $f$  and  $g$  are in non-parametric form, could also be wrong since it assumes an additive structure;
- Sometimes, a “wrong” model may work better than the true model for prediction purposes.

# Basic assumptions

---

- $X$  and  $Y$  have a joint distribution  $P_{X,Y}$ ;<sup>6</sup>
- The domain of  $Y$  and  $X$  are compact sets, respectively, i.e.,  $\mathcal{Y} := [-M_1, M_1]$  and  $\mathcal{X} := [-M_2, M_2]^d$ ;  $M_1$  and  $M_2$  are two positive constants.<sup>7</sup>

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We call our method **Model-free** since no restricted model format is assumed.

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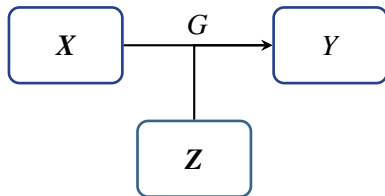
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Here,  $G : \mathcal{X} \times \mathcal{Z} \rightarrow \mathcal{Y}$ ;  $\mathcal{Z}$  is the domain of the reference random variable  $\mathbf{Z}$ .



## Noise outsourcing lemma

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**Lemma 1: Noise outsourcing** (Bloem-Reddy et al., 2020)

Let  $X$  and  $Y$  be random variables with joint distribution  $P_{X,Y}$ . Then, there is a measurable function  $G : [0, 1]^p \times \mathcal{X} \rightarrow \mathcal{Y}$  such that

$$Y \stackrel{a.s.}{=} G(X, Z), \text{ where } Z \sim \text{Uniform}[0, 1]^p \text{ and } Z \perp\!\!\!\perp X.$$

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In other words, the randomness in the conditional distribution of  $Y$  given  $X = \mathbf{x}$  is outsourced to reference random variable  $Z$  through  $G(\mathbf{x}, Z)$ .

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**Proposition 1:** A continuous counterpart of  $G(\cdot, \cdot)$  exists

Under some basic assumptions, we have a continuous  $\tilde{G}(\cdot, \cdot) : \mathcal{X} \times \mathcal{Z} \rightarrow \mathcal{Y}$  such that  $\tilde{G}(\mathbf{x}, z) = G(\mathbf{x}, z)$  for all  $(\mathbf{x}, z)$  except a negligible (in Lebesgue measure) set.

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To simplify the notation, we will keep using  $G(\cdot, \cdot)$  for this continuous counterpart. We will focus on estimating this continuous variant by Deep Neural Networks (DNN) and make predictions based on it.

# The structure of Deep Neural Networks

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In short, the structure of fully connected feedforward Deep Neural Networks (DNN) mainly depends on:

- The input and output dimensions;
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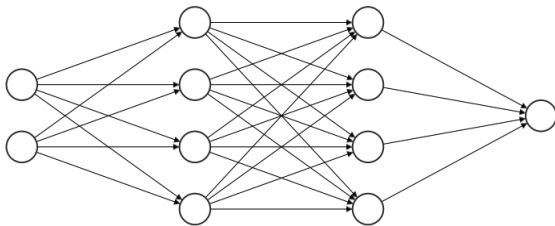
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We can write a DNN function  $f_{\text{DNN}}$  as

$$f_{\text{DNN}}(\mathbf{x}) = \mathbf{A}_{L+1}(\phi(\mathbf{A}_L(\cdots \phi(\mathbf{A}_3\phi(\mathbf{A}_2\phi(\mathbf{A}_1\mathbf{x} + \mathbf{b}_1) + \mathbf{b}_2) + \mathbf{b}_3) \cdots) + \mathbf{b}_L) + \mathbf{b}_{L+1};$$

where  $\{\mathbf{A}_i\}_{i=1}^{L+1}$  are weight matrices whose shapes depend on  $\mathbf{W}$  and output dimension;  $\{\mathbf{b}_i\}_{i=1}^{L+1}$  are intercept terms;  $\phi(\cdot)$  is the activation function.

**Figure 1:** The illustration of a fully connected DNN with  $L = 2$ ,  $W_1 = W_2 = 4$ ; input dimension and output dimension are 2 and 1, respectively.





# Training algorithm

---

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**Algorithm** Training procedure to get empirically optimal estimator  $\widehat{H}$

---

- 1: Initiate a DNN  $H_\theta \in \mathcal{F}_{\text{DNN}}$ <sup>8</sup> and simulate  $\{Z_i\}_{i=1}^n$  from  $P_Z$ .
- 2: **for** number of epochs **do**
- 3:     Update  $H_\theta$  by descending its gradient with the chosen optimization algorithm:

$$\nabla_\theta \left\{ \frac{1}{n} \sum_{i=1}^n (Y_i - H_\theta(\mathbf{X}_i, Z_i))^2 \right\}.$$

- 4:     Clip the parameter of  $H_\theta$  to  $[-m, m]$ .
  - 5: **end for**
  - 6: **Return** The estimated  $\widehat{H}(\cdot, \cdot)$ .
- 

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<sup>8</sup> $\mathcal{F}_{\text{DNN}}$  is a user-chosen space that contains all DNN candidates.

## Error bound for $\widehat{H}$

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**Theorem 4:** A high probability non-asymptotic error bound for  $\widehat{H}$

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When sample size  $n$  is large enough and under some further mild conditions, we have:

$$\left\| \widehat{H} - G \right\|_{L^2(X,Z)}^2 \leq C \cdot n^{-\frac{2}{\tau+d+p}} + o(n^{-\frac{2}{\tau+d+p}}); \text{ for } d + p \geq 2; \tau > 2;$$

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$$W := 3^{d+p+3} \max \left\{ (d+p) \left\lfloor N_1^{1/(d+p)} \right\rfloor, N_1 + 1 \right\}; \quad L := 12N_2 + 14 + 2(d+p); \quad N_1 = \left\lceil \frac{n^{\frac{d+p}{2(\tau+d+p)}}}{\log n} \right\rceil; \quad N_2 = \lceil \log(n) \rceil.$$



## Estimation of conditional distribution

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Define  $\widehat{F}_{\widehat{H}(\mathbf{x}_0, Z)}$  as the empirical distribution of  $\{\widehat{H}(\mathbf{x}_0, Z_i)\}_{i=1}^S$ ;  $S$  is the number of Monte Carlo sampling we apply to generate values of reference random variable.

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Under some additional restrictions about  $P_{X,Y}$ , we have

**Theorem 5:** Uniform estimation of  $F_{Y|X}$  based on  $\widehat{H}$

we have:

$$\sup_y \left| \widehat{F}_{\widehat{H}(\mathbf{x}_0, Z)}(y) - F_{Y|\mathbf{x}_0}(y) \right| \xrightarrow{P} 0, \text{ as } n \rightarrow \infty, S \rightarrow \infty,$$

for any  $\mathbf{x}_0 \in \mathcal{X}$  and  $y \in \mathcal{Y}$ , with probability at least  $1 - \exp(-n^{\frac{d+p}{\tau+d+p}})$ .

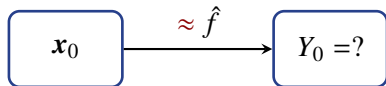
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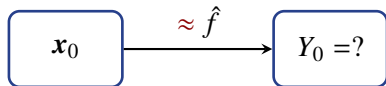
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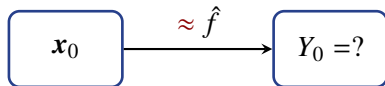
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An oracle  $G(\cdot, \cdot)$  can solve both error sources a.s.. However, error (2) still exists in practice.



# Preparations for PPI

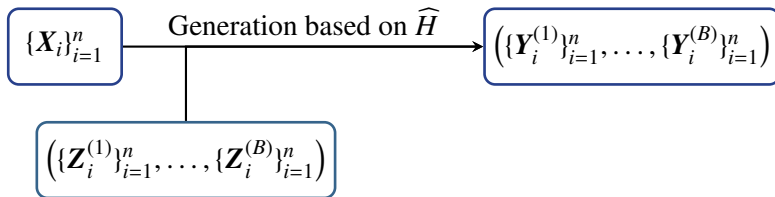
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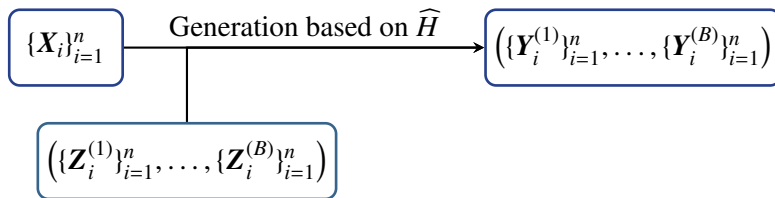
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Then, make re-estimation to get  $\{\widehat{H}^{(b)}\}_{b=1}^B$  based on  $(\{Y_i^{(1)}\}_{i=1}^n, \dots, \{Y_i^{(B)}\}_{i=1}^n)$ ,  $\{X_i\}_{i=1}^n$  and  $\{Z_i\}_{i=1}^n$ .

## The form of PPI based on $\hat{H}$

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Conditional on  $\{(X_i, Y_i, Z_i)\}_{i=1}^n$ , we approximate the predictive root  $R_0$  by the variant  $R_0^*$ :

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where,

- $R_0$  could be  $Y_0 - \widehat{Y}_{0,L_2}$ ;  $Y_0 \sim P_{Y|x_0}$  and  $\widehat{Y}_{0,L_2} := \mathbb{E}(\widehat{H}(\mathbf{x}_0, Z))$  is the *estimated* optimal  $L_2$  conditional point prediction; we approximate it by  $\frac{1}{S} \sum_{s=1}^S \widehat{H}(\mathbf{x}_0, Z_s)$ ;

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Thus, a pertinent PI with  $1 - \alpha$  coverage rate centered at  $\widehat{Y}_{0,L_2}$  has the form:

$$\left[ \widehat{Y}_{0,L_2} + Q_{\alpha/2}, \widehat{Y}_{0,L_2} + Q_{1-\alpha/2} \right];$$

$Q_{\alpha/2}$  and  $Q_{1-\alpha/2}$  are  $\alpha/2$  and  $1 - \alpha/2$  lower quantiles of  $P_{R_0^*}$ , the distribution of  $R_0^*$ . In practice,  $P_{R_0^*}$  can be approximated by the empirical distribution of  $\{Y_0^{(b)} - \widehat{Y}_{0,L_2}^{(b)}\}_{b=1}^B$ .

## Other DNN generative methods

Recently, Zhou et al. (2023) and Liu et al. (2021) proposed two conditional generators to estimate the conditional distribution in the regression context. Their methods rely on the adversarial training strategy which was first proposed by Goodfellow et al. (2014). We use  $\widehat{G}_{\text{KL}}$  and  $\widehat{G}_{\text{WA}}$  to represent these two DNN-based deep generators, they can be trained by the below formula:

$$(\widehat{G}_{\text{KL}}, \widehat{D}_{\text{KL}}) = \arg \min_{G_\rho \in \mathcal{F}'_{\text{DNN,G}}} \arg \max_{D_\phi \in \mathcal{F}'_{\text{DNN,D}}} \frac{1}{n} \sum_{i=1}^n D_\phi(G_\rho(Z_i, X_i), X_i) - \frac{1}{n} \sum_{i=1}^n \exp(D_\phi(Y_i, X_i));$$

$$(\widehat{G}_{\text{WA}}, \widehat{D}_{\text{WA}}) = \arg \min_{G_\rho \in \mathcal{F}_{\text{DNN,G}}} \arg \max_{D_\phi \in \mathcal{F}_{\text{DNN,D}}} \frac{1}{n} \sum_{i=1}^n D_\phi(G_\rho(Z_i, X_i), X_i) - \frac{1}{n} \sum_{i=1}^n D_\phi(Y_i, X_i).$$

- The objective functions are based on variants of KL-divergence and Wasserstein-1 distance;
- $D_\phi$  is the discriminator/critic trained together with generator  $G_\rho$  adversarially;
- $\mathcal{F}_{\cdot,\cdot}$  and  $\mathcal{F}'_{\cdot,\cdot}$  represent appropriate DNN classes.



## Simulation setting for PI

---

We take the below model to generate  $n$  training and  $T$  test data:

$$Y_i = X_{i,1}^2 + \exp(X_{i,2} + X_{i,3}/3) + X_{i,4} - X_{i,5} + (0.5 + X_{i,2}^2/2 + X_{i,5}^2/2) \cdot \varepsilon_i.$$

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To simplify notation, we denote all PIs by  $\widehat{\mathcal{I}}$  and we consider two coverage rates under different conditioning levels:

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We approximate  $\text{CV}_1$  by  $\frac{1}{T} \sum_{t=1}^T P(Y_0 \in \widehat{\mathcal{I}}|\mathbf{x}_t); \mathbf{x}_t$  is the  $t$ -th test point.





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We take same DNN structure for all methods.

We also consider the sample standard deviation  $\hat{\sigma}_{\text{PI}}$  of 2000  $\text{CV}_2$ . Besides, we present the average length (AL) of different PIs.

Simulation results

Table 1: Simulation results of  $CV_1$  with varying  $n$  and  $p$ .

	$CV_1$	AL	$CV_1$	AL	$CV_1$	AL
p = 5	n = 200		n = 500		n = 2000	
QPI	0.861(0.170)	5.487(1.054)	0.927(0.110)	6.734(1.463)	0.787(0.177)	3.621(0.855)
PPI	0.893(0.139)	6.208(1.384)	0.941(0.095)	7.258(1.808)	0.789(0.173)	3.728(0.959)
PI-KL	0.842(0.193)	5.496(0.861)	0.869(0.157)	5.434(1.218)	0.913(0.104)	5.670(2.282)
PI-WA	0.852(0.181)	5.439(0.907)	0.882(0.150)	5.970(2.030)	0.899(0.105)	5.365(1.996)
p = 10						
QPI	0.928(0.129)	7.497(0.720)	0.949(0.094)	8.194(0.950)	0.855(0.157)	4.474(0.817)
PPI	<b>0.944(0.105)</b>	8.103(1.072)	0.961(0.076)	8.623(1.325)	0.855(0.154)	4.546(0.953)
PI-KL	0.900(0.133)	6.701(0.835)	0.925(0.119)	6.806(0.933)	0.928(0.099)	5.882(1.403)
PI-WA	0.898(0.146)	6.757(0.719)	0.933(0.116)	7.545(1.340)	0.934(0.100)	6.199(1.880)
p = 15						
QPI	0.915(0.137)	7.408(0.669)	0.945(0.097)	7.430(0.949)	0.915(0.123)	5.895(0.647)
PPI	0.930(0.119)	7.760(0.936)	<b>0.953(0.085)</b>	7.749(1.172)	0.916(0.121)	5.971(0.807)
PI-KL	0.909(0.136)	7.427(0.817)	0.949(0.095)	8.082(1.068)	0.943(0.089)	6.556(1.491)
PI-WA	0.901(0.137)	6.797(0.687)	0.950(0.095)	7.972(1.312)	0.947(0.088)	6.778(1.541)
p = 20						
QPI	0.879(0.172)	6.726(0.485)	0.959(0.085)	8.830(0.683)	0.940(0.102)	6.849(0.562)
PPI	0.893(0.154)	6.941(0.702)	0.966(0.073)	9.100(0.950)	0.942(0.097)	6.925(0.759)
PI-KL	0.923(0.126)	7.799(0.842)	0.954(0.087)	8.311(0.861)	0.946(0.093)	6.806(1.097)
PI-WA	0.910(0.140)	7.402(0.698)	0.945(0.099)	8.011(0.800)	0.946(0.092)	6.804(1.534)
p = 25						
QPI	0.871(0.172)	7.020(0.287)	0.961(0.088)	9.633(0.645)	0.946(0.099)	7.296(0.475)
PPI	0.884(0.160)	7.189(0.548)	0.967(0.078)	9.881(0.938)	<b>0.948(0.095)</b>	7.370(0.695)
PI-KL	0.907(0.142)	7.370(0.618)	0.954(0.090)	8.670(0.813)	0.945(0.093)	6.915(1.009)
PI-WA	0.897(0.151)	7.071(0.510)	0.960(0.081)	8.514(0.942)	0.944(0.097)	7.117(1.491)

## Simulation results of $CV_2$ : PPI vs PI-KL

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# Simulation results of $CV_2$ : PPI vs PI-KL

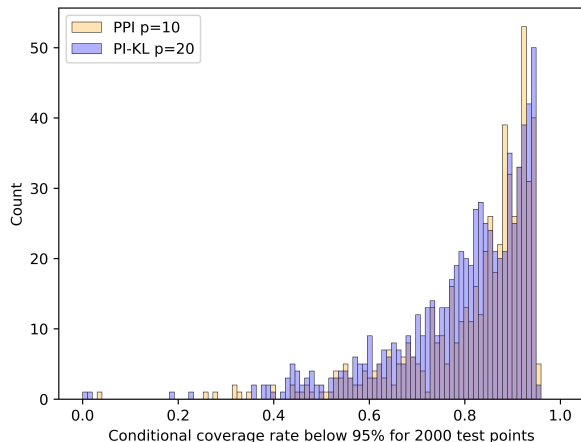
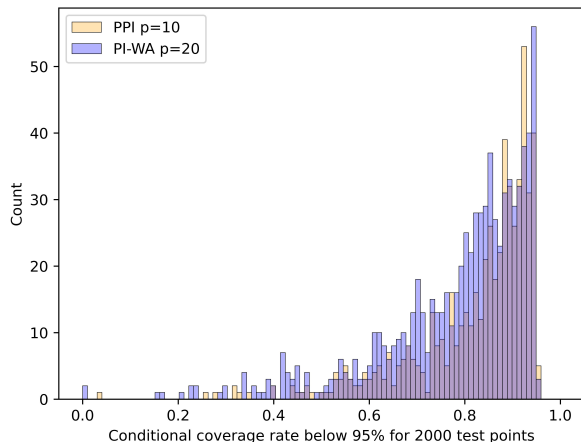


Figure 2: Histograms of all undercoverage  $CV_2$  ( $CV_2$  less than nominal level 95%) of PPI and PI-KL.

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**Figure 3:** Histograms of all undercoverage  $CV_2$  ( $CV_2$  less than nominal level 95%) of PPI and PI-WA.

## Simulation results of $CV_2$ : PPI vs QPI

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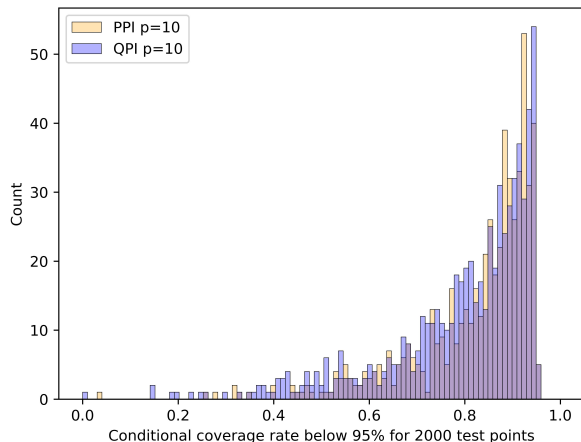


Figure 4: Histograms of all undercoverage  $CV_2$  ( $CV_2$  less than nominal level 95%) of PPI and QPI.

# Theoretical explanations

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Under further assumptions about the joint distribution  $P_{X,Y}$ , we have:

## Theorem 6: Theoretical understanding of PPI with DNN

For an appropriate sequence of sets  $\Omega_n$ , such that  $\mathbb{P}(\{X_i, Y_i, Z_i\}_{i=1}^n \notin \Omega_n) = o(1)$ , PPI can capture the estimation variability under  $S \rightarrow \infty$  in an appropriate rate for each  $n$ , when  $n \rightarrow \infty$ .

Furthermore,

$$\sup_y \left| \widehat{F}_{\widehat{H}(\mathbf{x}_0, Z)} \star \phi_\sigma(y) - F_{Y|\mathbf{x}_0} \star \phi_\sigma(y) \right| \leq \sup_y \left| \widehat{F}_{\widehat{H}(\mathbf{x}_0, Z)}(y) - F_{Y|\mathbf{x}_0}(y) \right| \text{ with probability } 1;$$

$\widehat{F}_{\widehat{H}(\mathbf{x}_0, Z)}$  is the empirical distribution of  $\{\widehat{H}(\mathbf{x}_0, Z_i)\}_{i=1}^S$ ;  $\star$  is the convolution operator;  $\phi_\sigma$  is the density function of the normal distribution  $N(0, \sigma^2)$ .

## Scalable Subsampling for Deep Neural Networks training<sup>9</sup>

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<sup>9</sup>This part is based on:

- Wu, K. and Politis, D.N., Scalable Subsampling Inference of Deep Neural Networks. *ACM/IMS Journal of Data Science* 2025, 2(1), 1–29.

# Motivation

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From the part II, we have seen the power of DNN. However, there are two crucial problems that need more thoughts:



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- **The computational burden of training DNN:** DNN has been developed rapidly fueled by ever-increasing amounts of data. However, training a large DNN with a huge sample size requires heavy computational resources.

We want to kill two birds with one stone.



- **Variance reduction:** To improve the convergence rate of an estimator, we can try to decrease its variance if its bias is acceptable. This is inspired by *bagging* method of Breiman (1996).

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# Intuition

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## Example: A simple implementation

Suppose we need  $O(n^\zeta)$  operations to compute one estimator  $\hat{\theta}_n$  for true parameter  $\theta$ . If we consider  $q = O(n/b)$  number of estimations  $\hat{\theta}_{b,i}$  on subsamples with size  $b$  for  $i = 1, \dots, q$ , we can take  $\bar{\theta}_{b,n,ss} := \frac{1}{q} \sum_{i=1}^q \hat{\theta}_{b,i}$  to approximate  $\hat{\theta}_n$ . Then, only  $O(nb^{\zeta-1})$  operations are needed.



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Assume that we observe the sample  $\{U_1, \dots, U_n\}$ ;  $U_i$  represents  $(X_i, Y_i)$ ; then, scalable subsampling relies on  $q = \lfloor (n - b)/h \rfloor + 1$  number of subsamples  $B_1, \dots, B_q$  where  $B_j = \{U_{(j-1)h+1}, \dots, U_{(j-1)h+b}\}$ ; here,  $\lfloor \cdot \rfloor$  denotes the floor function, and  $h$  controls the amount of overlap (or separation) between  $B_j$  and  $B_{j+1}$ .

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# Scalable subsampling

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Recently, Politis (2024) proposed one type of non-stochastic *scalable subsampling* technique.

Assume that we observe the sample  $\{U_1, \dots, U_n\}$ ;  $U_i$  represents  $(X_i, Y_i)$ ; then, scalable subsampling relies on  $q = \lfloor (n - b)/h \rfloor + 1$  number of subsamples  $B_1, \dots, B_q$  where  $B_j = \{U_{(j-1)h+1}, \dots, U_{(j-1)h+b}\}$ ; here,  $\lfloor \cdot \rfloor$  denotes the floor function, and  $h$  controls the amount of overlap (or separation) between  $B_j$  and  $B_{j+1}$ .

Tuning  $b$  and  $h$  can make scalable subsampling samples have different overlapping rates:

- if  $h = 1$ , the overlap is the maximum possible;
- if  $h = 0.2b$ , there is 80% overlap between  $B_j$  and  $B_{j+1}$ ;
- if  $h = b$ , there is no overlap between  $B_j$  and  $B_{j+1}$  but these two blocks are adjacent;
- if  $h = 1.2b$ , there is a block of about  $0.2b$  data points that separate the blocks  $B_j$  and  $B_{j+1}$ .

## Scalable subsampling estimation of DNN

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Define the scalable subsampling DNN estimator as  $\bar{f}_{\text{DNN}}(\mathbf{X}) = \frac{1}{q} \sum_{j=1}^q \widehat{f}_{\text{DNN},b,j}(\mathbf{X})$ ; here,  $q = \lfloor (n - b)/h \rfloor + 1$ , and  $\widehat{f}_{\text{DNN},b,j}(\cdot)$  is trained DNN with the  $j$ -th subsample  $B_j$ .

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Under below assumptions:

- $\mathbb{E}(\widehat{f}_{\text{DNN}}(\mathbf{x}) - f(\mathbf{x})) = O(n^{-\Lambda/2})$  uniformly for all  $\mathbf{x} \in \mathcal{X}$  for some constant  $\Lambda > 0$ ;

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we have

## Theorem 7:

Under appropriate assumptions, let  $b = h = n^\beta$ ;  $\beta = \frac{1}{1+\Lambda-\frac{\xi}{\xi+d}}$ . Then, with probability at least  $(1 - \exp(-n^{\frac{d}{\xi+d}} \log^6 n))^q$ :

$$\left\| \bar{f}_{\text{DNN}} - f \right\|_{L_2(X)}^2 \leq n^{\frac{-\Lambda}{\Lambda+\frac{d}{\xi+d}}} \mathcal{L}(n);$$

where  $\mathcal{L}(n)$  is a slowly varying function involving a constant and all  $\log(n)$  terms.

## Simulation setting

---

To perform simulations, we consider below models:

- Model-1:  $Y_i = X_{i,1}^2 + \sin(X_{i,2} + X_{i,3}) + \epsilon$ , where  $\mathbf{X} \sim N(0, \mathbf{I}_3)$ ;  $\epsilon \sim N(0, 1)$ ;
- Model-2:  $Y_i = X_{i,1}^2 + \sin(X_{i,2} + X_{i,3}) + \exp(-|X_{i,4} + X_{i,5}|) + \epsilon$ , where  $\mathbf{X} \sim N(0, \mathbf{I}_5)$ ;  $\epsilon \sim N(0, 1)$ .

To be consistent with folk wisdom, we build  $\widehat{f}_{\text{DNN},b,i}$  with a relatively large depth to decrease the bias but also guarantee that the DNN estimator is in the under-parameterized region. We also consider other 5 DNN estimators trained with the whole sample:

- (1) A DNN possesses the same depth and width as  $\widehat{f}_{\text{DNN},b,i}$ . We denote it “S-DNN”;
- (2) A DNN possesses the same depth as  $\widehat{f}_{\text{DNN},b,i}$ , but a larger width so that its parameter size is close to the sample size. We denote it “DNN-deep-1”;
- (3) A DNN possesses the same depth as  $\widehat{f}_{\text{DNN},b,i}$ , but a larger width so that its parameter size is close to half of the sample size. We denote it “DNN-deep-2”;
- (4) A DNN possesses only one hidden layer, but a larger width so that its parameter size is close to the sample size. We denote it “DNN-wide-1”;
- (5) A DNN possesses only one hidden layer, but a larger width so that its parameter size is close to half of the sample size. We denote it “DNN-wide-2”.



# Hyperparameter setting and metric

---

To evaluate the performance of different DNN estimators, we apply the empirical MSE and MSPE criteria:

$$\text{MSE} := \frac{1}{n} \sum_{i=1}^n (\tilde{f}_{\text{DNN}}(\mathbf{x}_i) - f(\mathbf{x}_i))^2 ; \text{MSPE} := \frac{1}{N} \sum_{i=1}^N (\tilde{f}_{\text{DNN}}(\mathbf{x}_{0,i}) - f(\mathbf{x}_{0,i}))^2 ;$$

here  $\tilde{f}_{\text{DNN}}(\cdot)$  represents different DNN estimators and  $f(\cdot)$  is the true regression function;  $\{\mathbf{x}_i, y_i\}_{i=1}^n$  are training data;  $\{\mathbf{x}_{0,i}, y_{0,i}\}_{i=1}^N$  are test data; we take  $N = 2 \cdot 10^5$ .

Simulation results are averaged from 200 replications.



# Simulation results

Table 2: MSE/MSPE and Training Time (in seconds) of different DNN models

Estimator:	SS-DNN	S-DNN	DNN-deep-1	DNN-deep-2	DNN-wide-1	DNN-wide-2
Model-1, $n = 10^4$						
Width	[15,15,15]	[15,15,15]	[65,65,65]	[45,45,45]	[2000]	[1000]
MSE	0.0296	0.0536	0.0533	0.0522	0.0426	0.0431
MSPE	0.0310	0.0564	0.0572	0.0570	0.0453	0.0449
Training Time	353	379	561	468	483	363
Model-2, $n = 10^4$						
Width	[15,15,15]	[15,15,15]	[65,65,65]	[45,45,45]	[2000]	[1000]
MSE	0.0757	0.0830	0.1076	0.0980	0.0729	0.0728
MSPE	0.0790	0.0875	0.1114	0.1045	0.0754	0.0749
Training Time	359	376	560	471	551	394
Model-2, $n = 2 \cdot 10^4$						
Width	[20,20,20]	[20,20,20]	[95,95,95]	[65,65,65]	[2800]	[1400]
MSE	0.0490	0.0653	0.0686	0.0675	0.0635	0.0635
MSPE	0.0502	0.0670	0.0692	0.0689	0.0623	0.0626
Training Time	748	775	1684	1198	1549	998

# Acknowledgements I

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I would like to thank Professor Dimitris Politis. As my advisor, my friend, and my life guider, he always supported my research and gave me valuable suggestions. Moreover, he led me to a path I never imagined about.

## Acknowledgements II

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I would like to thank my committee members, Ery Arias-Castro, Yian Ma and Danna Zhang.

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# Acknowledgements III

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I would like to thank all my friends I met during my Ph.D. life.



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I would also like to thank the administrative staff, particularly Scott Rollans and Mark Whelan, for their constant support.

# Acknowledgements IV

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I would like to thank my parents. This talk is dedicated to them.

Thank you!

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## **Backup I: Additional slides**

# Estimation of $G(\cdot, \cdot)$

---

Define the optimal estimator:

$$H_0 = \arg \min_H \mathbb{E} (Y - H(X, Z))^2 = \arg \min_H \mathcal{R};$$

A simple decomposition shows that:

$$\mathbb{E} (Y - H(X, Z))^2 = \mathbb{E} (Y - G(X, Z) + G(X, Z) - H(X, Z))^2 .$$

Thus, the optimal  $H_0$  is unique almost surely and we can take it as the continuous counterpart of  $G$ .

# Difference between traditional MSE risk

Recall that the risk for standard regression tasks is

$$\mathbb{E}[(Y - h(X))^2] := \mathcal{R}_s.$$

Table 3: Comparison between standard regression risk and our risk

	Geometry	$\sigma$ -algebra
$\mathcal{R}_s$	The optimal estimator is the projection of $Y$ onto a closed subspace $\mathcal{S}_X$ of $L_2$ consisting of all random variables which can be written in a function of $X$ .	$\mathbb{E}(Y X)$ is $\mathcal{D}_X$ -measurable. <sup>10</sup>
$\mathcal{R}$	The optimal estimator is a projection of $Y$ onto an extended version of $\mathcal{S}_X$ by random variable $Z$ .	$Y \stackrel{a.s.}{=} G(X, Z)$ is $\mathcal{D}_{(X,Z)}$ -measurable.

<sup>10</sup> $\mathcal{D}_X$  is the  $\sigma$ -algebra generated by  $X$ ;  $\mathbb{E}(Y|X)$  could also equal to  $Y$  a.s. if  $Y$  is  $\mathcal{D}_X$ -measurable, e.g.,  $\mathbb{E}(Y|Y) = Y$ .



# Intuition behind our Deep limit model-free prediction algorithm

We provide a toy example to explain the motivation of our training procedure.

## Remark: An illustration example

Suppose we need to estimate the coefficient  $\beta$  of a linear regression model  $Y = \beta^T \cdot X + \epsilon$  with a fixed design based on samples  $\{(\mathbf{x}_i, y_i)\}_{i=1}^n$ ; here,  $\epsilon$  has zero mean and finite variance.

- OLS:  $\widehat{\beta} := \arg \min_{\beta} \frac{1}{n} \sum_{i=1}^n (y_i - \beta_i^T \cdot \mathbf{x}_i)^2$  which is consistent under standard conditions.
- Variant of OLS:  $\widehat{\beta}^* := \arg \min_{\beta} \frac{1}{n} \sum_{i=1}^n (y_i - (\beta_i^T \cdot \mathbf{x}_i + \epsilon_i^*))^2$  where  $\{\epsilon_i^*\}_{i=1}^n$  are independent of  $X$  and can be generated from any distribution with mean zero and finite variance.

$\widehat{\beta}^*$  is also consistent although  $\widehat{\beta}$  would generally be more efficient.

Analogously, our DNN-based estimation  $\widehat{H}^*$  converges to  $H_0$  in the mean square sense even using the artificially generated  $\{Z_i^*\}_{i=1}^n$ .

The estimation error of  $\widehat{H}$  can be decomposed into two sources:

- (1) The stochastic error, which measures the difference between  $\widehat{H}$  and the best estimator  $H^*$  in a DNN class  $\mathcal{F}_{\text{DNN}}$ ;  
$$H^* := \arg \min_{H \in \mathcal{F}_{\text{DNN}}} \|H_0 - H\|_{\infty};$$
- (2) The approximation error, which measures the difference between  $H_0$  and  $H^*$  in a DNN class  $\mathcal{F}_{\text{DNN}}$ .

Table 4: Comparison between different DNN-based methods

$\widehat{H}$		$\widehat{G}_{\text{KL}}, \widehat{G}_{\text{WA}}$
Stability	The training process is more stable and directly due to the MSE-like loss function.	The training process is sensitive to the training setting and depends on $D_\phi$ being optimal given current step $G_\rho$ .
Metrics	The optimization corresponds to minimizing the Kolmogorov distance between two distributions.	The optimization corresponds to minimizing KL-divergence and Wasserstein-1 distance <sup>11</sup> .
Computability	Only one DNN need to be trained.	Two DNNs need to be trained adversarially.

<sup>11</sup>The “distance” between two distributions converges to 0 under the metric of Wasserstein-1 distance or KL-divergence implies the convergence measured by Kolmogorov distance.

## Simulation setting for optimal $L_2$ point prediction

---

We take the below model from Zhou et al. (2023) to generate  $n$  training and  $T$  test data:

$$Y_i = X_{i,1}^2 + \exp(X_{i,2} + X_{i,3}/3) + X_{i,4} - X_{i,5} + (0.5 + X_{i,2}^2/2 + X_{i,5}^2/2) \cdot \varepsilon_i;$$

where  $X_i$  and  $\varepsilon_i$  come from  $N(0, I_5)$  and  $N(0, 1)$  truncated to  $[-5, 5]^5$  and  $[-5, 5]$ , respectively.

To predict the mean of  $Y$  conditional on  $X = \mathbf{x}$ , we rely on  $\widehat{Y}_t = \frac{1}{S} \sum_{s=1}^S \widehat{\Pi}(\mathbf{x}_t, Z_s)$ ;  $Z_s \sim N(0, I_p)$ ;  $\mathbf{x}_t$  is the  $t$ -th observation of the test data;  $\widehat{\Pi}$  represents trained model  $\widehat{H}$ ,  $\widehat{G}_{\text{KL}}$  or  $\widehat{G}_{\text{WA}}$ .

To measure different methods, we repeat the simulations  $K$  times and consider the metric:

$$\widetilde{L} = \frac{1}{T} \sum_{t=1}^T \frac{1}{K} \sum_{k=1}^K (Y_{t,L_2} - \widehat{Y}_{k,t})^2;$$

where  $Y_{t,L_2}$  is the oracle  $L_2$  optimal value of  $Y$  conditional on  $\mathbf{x}_t$ ;  $\widehat{Y}_{k,t}$  is the conditional  $L_2$  point prediction based on the  $k$ -th training data.

# Simulation setting

---

We apply the same hyperparameter setting to train all DNN.

We take  $n = 2000$ ,  $T = 2000$ ,  $S = 10000$ ,  $K = 200$  to compute the error metric.

For the structure of DNN, we separate the simulation studies into two groups: (a) structures of  $\widehat{H}$  and  $\widehat{G}$  are [35,35] and [50]<sup>12</sup>, respectively; (b) structures of  $\widehat{H}$  and  $\widehat{G}$  are all [35,35]. For both groups,  $\widehat{D}$  takes the same structure as the previous work, i.e., [50,25].

For the benchmark method, we apply the numerical integration  $\int_{\mathcal{Y}} y \hat{f}_{y|\mathbf{x}_t} dy$  with 1000 subdivisions to approximate  $E(Y|\mathbf{x}_t)$ ;  $\hat{f}_{y|\mathbf{x}_t}$  is the kernel conditional density estimator of  $Y$  conditional on  $\mathbf{x}_t$ .

---

<sup>12</sup>[35,35] stands for a two layers DNN and each layer has 35 neurons; [50] is the DNN structure used in Zhou et al. (2023). To simplify notations,  $\widehat{G}$  represents  $\widehat{G}_{\text{KL}}$  or  $\widehat{G}_{\text{WA}}$ ;  $\widehat{D}$  represents  $\widehat{D}_{\text{KL}}$  or  $\widehat{D}_{\text{WA}}$ .

Table 5: Point predictions of different methods under groups (a) and (b).

	Group (a)			Group (b)		
	$\widehat{H}$	$\widehat{G}_{\text{KL}}$	$\widehat{G}_{\text{WA}}$	$\widehat{H}$	$\widehat{G}_{\text{KL}}$	$\widehat{G}_{\text{WA}}$
SGD						
$p = 1$	0.309	3.931	10.39	0.292	3.827	82.97
$p = 3$	0.298	4.009	11.10	0.285	3.762	56644
$p = 5$	0.296	4.036	40.39	0.281	3.801	12843
$p = 10$	<b>0.294</b>	4.116	182.3	<b>0.280</b>	3.812	11378
Adam						
$p = 1$	1.608	1.838	3558	1.572	1.836	14322
$p = 3$	0.832	1.105	8.480	0.843	1.549	43.48
$p = 5$	0.604	0.820	43.85	0.591	1.166	43.84
$p = 10$	<b>0.412</b>	0.495	5.523	<b>0.422</b>	0.817	14.50
RMSProp						
$p = 1$	0.960	1.767	1.910	0.973	1.620	2.326
$p = 3$	0.601	1.049	1.248	0.597	0.964	1.263
$p = 5$	0.484	0.779	0.908	0.479	0.727	0.903
$p = 10$	<b>0.365</b>	0.463	0.598	<b>0.352</b>	0.494	0.508

Note: The error metric  $\widetilde{L}$  of using conditional kernel density estimation is around 1.210.

# Hyperparameter setting

---

We apply the same hyperparameter setting to train  $\widehat{H}$ ,  $\widehat{G}_{\text{KL}}$  and  $\widehat{G}_{\text{WA}}$ :  $n = 2000$ ;  $T = 2000$ ;  $S = 10000$ ;  $K = 200$ ;  $p = 1, 3, 5, 10$ ,  $m = 20$ ; Learning rate: 0.001; Number of epochs: 10000.

For the optimizer of the adversarial training process, Arjovsky et al. (2017) proposed using optimizer RMSProp with Wasserstein distance is more appropriate. However, Pang et al. (2020) argued that SGD-based optimizers are better. We consider three common optimizers, SGD, Adam and RMSProp.

## Remark of Theorem 3

---

- **PPI can capture the estimation variability:** Since the distribution of  $R_0^*$  can approximate the distribution of  $R_0$ , PPI captures the estimation variability in finite sample cases to some extent.
- **A convolution implied in predictive root:** It comes from rewriting the predictive root as  $R_0 := Y_0 - \mathbb{E}(Y_0|\mathbf{x}_0) + \mathbb{E}(Y_0|\mathbf{x}_0) - \widehat{Y}_{0,L_2}$ ;  $Y_0 - \mathbb{E}(Y_0|\mathbf{x}_0)$  only depends on  $P_{Y|\mathbf{x}_0}$  and  $\mathbb{E}(Y_0|\mathbf{x}_0) - \widehat{Y}_{0,L_2}$  is a (asymptotically shrinking) Gaussian distribution. Thus the below inequality from the previous theorem reveals that we need less data to achieve the same accuracy of the distribution estimation under this convolution approach.

$$\sup_y \left| \widehat{F}_{\widehat{H}(x_0, Z)} \star \phi_\sigma(y) - F_{Y|\mathbf{x}_0} \star \phi_\sigma(y) \right| \leq \sup_y \left| \widehat{F}_{\widehat{H}(x_0, Z)}(y) - F_{Y|\mathbf{x}_0}(y) \right| \text{ with probability 1.}$$



# KL-divergence and Wasserstein-1 distance

---

- KL-divergence: if  $f, g$  are densities of the measures  $\mu, \nu$  with respect to a dominating measure  $\lambda$ ,

$$d_I(\mu, \nu) := \int_{S(\mu)} f \log(f/g) d\lambda.$$

where  $S(\mu)$  is the support of  $\mu$  on  $\Omega$ .

- Wasserstein-1 distance: for  $\Omega = \mathbb{R}$ , if  $F, G$  are the distribution functions of  $\mu, \nu$  respectively, the Kantorovich metric is defined by

$$\begin{aligned} d_W(\mu, \nu) &:= \int_{-\infty}^{\infty} |F(x) - G(x)| dx \\ &= \int_0^1 |F^{-1}(t) - G^{-1}(t)| dt. \end{aligned}$$

## Remark on scalable subsampling method

---

The scalable subsampling method can be applied in making point estimations and developing the estimation inference:

- **For point estimation:** Take  $h = b$  as an example, as the analysis in the previous example,  $O(nb^{\zeta-1})$  operations are needed to compute  $\bar{\theta}_{b,n,SS}$ . Moreover, we have

$$\mathbb{E}(\bar{\theta}_{b,n,SS}) = \mathbb{E}(\hat{\theta}_{b,1}) \text{ and } \text{Var}(\bar{\theta}_{b,n,SS}) \leq \frac{1}{q} \text{Var}(\hat{\theta}_{b,1}); q = \lfloor n/h \rfloor.$$

Hence, if the bias of  $\hat{\theta}_{b,1}$  is tolerable,  $\bar{\theta}_{b,n,SS}$  yields a welcome variance reduction.

- **For estimation inference:** The subsampling distribution  $L_{n,b,h}(x) = q^{-1} \sum_{i=1}^q \mathbb{1} \{ \tau_b g(\hat{\theta}_{b,i} - \hat{\theta}_n) \leq x \}$  can be used to approximate the distribution of the estimation root  $J_n(x) = P \{ \tau_n g(\hat{\theta}_n - \theta) \leq x \}$  under mild conditions; where  $g(\cdot)$  could be the identity function or the sup-norm.

## Remark on scalable subsampling estimation of DNN

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  - (2) As revealed in Belkin et al. (2019), the double-descent of the risk exists for over-parameterized estimator. Thus, we may take  $\Delta > n$  to meet the bias order requirement.

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- The two assumptions regarding the bias order of the DNN estimator are feasible:
  - (1) As revealed in Yarotsky and Zhevnerchuk (2020), the approximation ability in the uniform sup-norm of a DNN can be as fast as  $\Delta^{-2\xi/d}$ ;  $\Delta$  is the total number of parameters in a DNN. Although this rate is not instructive in practice, the desired bias order is achievable.
  - (2) As revealed in Belkin et al. (2019), the double-descent of the risk exists for over-parameterized estimator. Thus, we may take  $\Delta > n$  to meet the bias order requirement.
- The saving of computational cost from applying scalable subagging will be more significant for executing estimation with a large DNN or with a large sample. Assume that a DNN with size  $W = \Theta(n^\phi)$ . The total number of operations to train a DNN is  $O(n \cdot W \cdot E)$ ; here  $E$  represents the number of epochs. When the size of the DNN is larger than the sample size,  $O(n \cdot W \cdot E) \approx O(n^\varphi)$ ;  $\varphi > 2$ . For our estimator, the number of operations is  $O(n^{\beta\varphi} q) = O(n^{1+\beta(\varphi-1)})$ . The ratio of  $n^{1+\beta(\varphi-1)}$  over  $n^\varphi$  is  $n^{-(\varphi-1)(1-\beta)}$ .