

Significance tests for feature relevance of a black-box learner

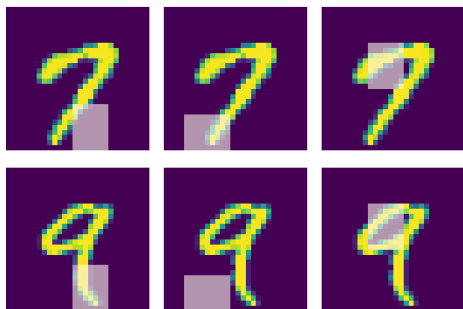
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<https://arxiv.org/abs/2103.04985>

<https://dnn-inference.readthedocs.io>

Illustrative example



- **Question:** Can we provide a **valid p -value** for **any pre-specified region** (features) based on a **black-box model**, such as a convolutional neural network?

Motivation

- **Why significance tests?** Hypothesis testing, feature interpretation, XAI, make black-box models more reliable ...
- **Why region tests?** For image analysis, the impact of each pixel is negligible but a pattern of a collection of pixels (e.g. a region) may instead become salient.
- **Why black-box models?** Significant improvement in prediction performance, which enforce us to believe that a black-box model is a better option to model real data. For example, use a CNN to formulate image data.

Difficulty

- **Black-box models.** It is infeasible (or difficult) to “open the box”, that is, we only access the input and output for a black-box model, and do not know its inner structure.
- **Feature-param correspondence.** The feature-parameter correspondence is unclear for black-box models, such as CNNs and RNNs.
- **High-dimensional hypothesized features.** The dimension of the hypothesized features could be extremely large.
- **Computationally expensive.** Refitting a deep learning model is computationally expensive.

Difficulty

- **Overparametrized models.** When the number of parameters increase, both training / testing errors decrease, and the training error could be zero.

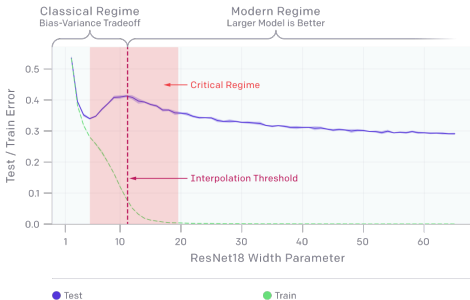


Figure 1: Source¹

¹<https://openai.com/blog/deep-double-descent/>

Issues for existing methods

- Likelihood Ratio Test (LRT)
 - **black-box models and overparam**: Taylor expansion is infeasible, and the training loss could be very small.
 - **feature-param relation** is unclear: LRT works for $\theta \in \Theta$ vs. $\theta \notin \Theta$.

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- Conditional randomization test (CRT; [Candès et al., 2018]) and holdout randomization test (HRT; [Tansey et al., 2018])
 - significance testing for a single feature.
 - require conditional Prob of hypothesized feature given the others. It is usually difficult to estimate for real complex datasets, or **high-dimensional hypothesized features**.

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 - require conditional Prob of hypothesized feature given the others. It is usually difficult to estimate for real complex datasets, or **high-dimensional hypothesized features**.
- Leave-one-covariate-out (LOCO; [Lei et al., 2018])
 - significance testing for a single feature.
 - finite-sample hypothesis testing.

Goodness

- Input and output: $\mathbf{X} \in \mathbb{R}^d$ and $\mathbf{Y} \in \mathbb{R}^K$;
 - **large-scale dataset** $(\mathbf{X}_i, \mathbf{Y}_i)_{i=1}^N$
- Black-box model: $f : \mathbb{R}^d \rightarrow \mathbb{R}^K$;
 - **good performance, or small generalization error, or reasonable convergence rate**
- Flexible computing platform for a general loss function $l(f(\mathbf{X}), \mathbf{Y})$
 - **TensorFlow, Keras, Pytorch**

The proposed risk-based testing

- **Goal:** testing the relevance of a sub-feature $\mathbf{X}_{\mathcal{S}} = \{X_j : j \in \mathcal{S}\}$ to the outcome \mathbf{Y} without specifying any form of the prediction function, where \mathcal{S} is an index set of hypothesized features.
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- **Our testing:** directly compare perf w/- or w/o hypothesized features
 - Masked data (\mathbf{Z}, \mathbf{Y}) , with permutation of $\mathbf{Z}_{\mathcal{S}}$ or $\mathbf{Z}_{\mathcal{S}} = \mathbf{0}$, and $\mathbf{Z}_{\mathcal{S}^c} = \mathbf{X}_{\mathcal{S}^c}$.
 - Risk functions:

$$R(f) = \mathbb{E}(l(f(\mathbf{X}), \mathbf{Y})), \quad R_{\mathcal{S}}(g) = \mathbb{E}(l(g(\mathbf{Z}), \mathbf{Y}))$$

- Population minimizer:

$$f^* = \underset{f}{\operatorname{argmin}} R(f), \quad g^* = \underset{g}{\operatorname{argmin}} R_{\mathcal{S}}(g)$$

$$H_0 : R(f^*) - R_{\mathcal{S}}(g^*) = 0, \quad \text{versus} \quad H_a : R(f^*) - R_{\mathcal{S}}(g^*) < 0. \quad (1)$$

The proposed risk-based testing

Relationships among the risk invariance hypothesis in (2), marginal independence, and conditional independence; the latter two are defined as:

Marginal indep: $\mathbf{Y} \perp \mathbf{X}_S$, conditional indep : $\mathbf{Y} \perp \mathbf{X}_S \mid \mathbf{X}_{S^c}$.

Lemma 1 (Relation to independence)

For any loss function, conditional independent implies risk invariance, or

$$\mathbf{Y} \perp \mathbf{X}_S \mid \mathbf{X}_{S^c} \implies R(f^*) - R_S(g^*) = 0.$$

Moreover, if the cross-entropy loss $l(f(\mathbf{X}), Y) = -\mathbf{1}_Y^\top \log(f(\mathbf{X}))$ is used in (2), then H_0 is equivalent to conditional independence almost surely under the marginal distribution of \mathbf{X} .

The proposed risk-based testing

- (Constant loss): $I(f(\mathbf{X}), Y) = C$.
- (L_2 -loss): $I(f(\mathbf{X}), Y) = \mathbb{E}(Y - f(\mathbf{X}))^2$.
- (Cross-entropy loss): $I(f(\mathbf{X}), Y) = \mathbf{1}_Y^T \log(f(\mathbf{X}))$.

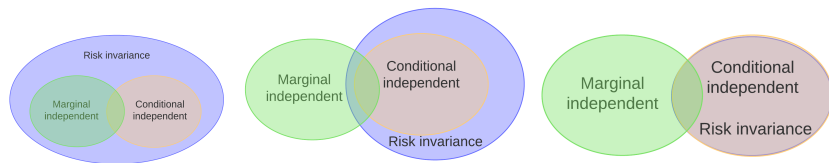
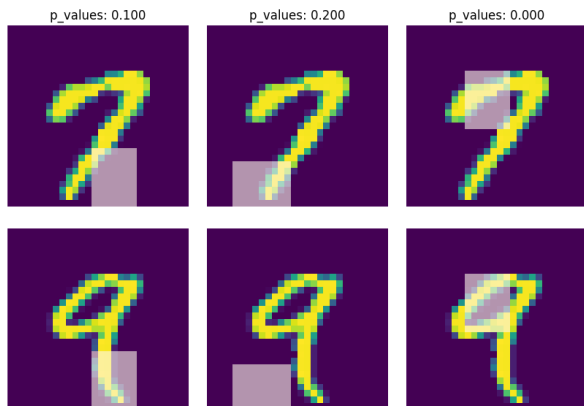


Figure 2: Three cases illustrate relationships among marginal independence, conditional independence, and risk invariance.

Our solution

- The proposed test is able to produce a valid p -value for (2).
- Python library `dnn-inference`
(<https://dnn-inference.readthedocs.io>)



Splitting data

- Recall the proposed hypothesis:

$$H_0 : R(f^*) - R_S(g^*) = 0, \quad \text{versus} \quad H_a : R(f^*) - R_S(g^*) < 0. \quad (2)$$

- Empirically** { estimate (f^*, g^*) , evaluate (R, R_S) }

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Question: do we need to split data? **Yes!**

- Deep neural networks easily fit shuffled pixels, random pixels.* See Figure 1 in [Zhang et al., 2016]: training loss converge to zero under random pixels, yet the testing loss is still sensible.
- Theoretically, it is not easy to find a limiting distribution based on a black-box model.

Splitting data

- Splitting data into **estimation** and **inference** sets

Total set $(\mathbf{X}_i, \mathbf{Y}_i)_{i=1}^N \rightarrow$ Est set $(\mathbf{X}_i, \mathbf{Y}_i)_{i=1}^n +$ Inf set $(\mathbf{X}_{n+j}, \mathbf{Y}_{n+j})_{j=1}^m$
 $(\mathbf{Z}_i, \mathbf{Y}_i)_{i=1}^n$ $(\mathbf{Z}_{n+j}, \mathbf{Y}_{n+j})_{j=1}^m$

- Obtain estimator (\hat{f}_n, \hat{g}_n) based on estimation set, then plug into evaluation on an inference sample:

$$\hat{R}(\hat{f}_n) - \hat{R}_S(\hat{g}_n)$$

- **Question:** Is it good estimation of $R(f^*) - R_S(g^*)$? Asymptotic null distribution?

Decomposition

- Compare $\widehat{R}(\widehat{f}_n) - \widehat{R}_S(\widehat{g}_n)$ with $R(f^*) - R_S(g^*)$
- Consider the following decomposition

$$\begin{aligned}\widehat{R}(\widehat{f}_n) - \widehat{R}_S(\widehat{g}_n) &= \widehat{R}(\widehat{f}_n) - R(\widehat{f}_n) + R_S(\widehat{g}_n) - \widehat{R}_S(\widehat{g}_n) \\ &\quad + R(\widehat{f}_n) - R(f^*) + R_S(g^*) - R_S(\widehat{g}_n) \\ &\quad + R(f^*) - R_S(g^*) = T_1 + T_2 + T_3\end{aligned}$$

- T_1 is a conditional IID sum

$$\begin{aligned}T_1 &= \widehat{R}(\widehat{f}_n) - R(\widehat{f}_n) + R_S(\widehat{g}_n) - \widehat{R}_S(\widehat{g}_n) \\ &= \frac{1}{m} \sum_{j=1}^m \left(\Delta_{n,j} - \mathbb{E}(\Delta_{n,j} | (\mathbf{X}_i, \mathbf{Y}_i)_{i=1}^n) \right),\end{aligned}$$

where $\Delta_{n,j} = l(\widehat{f}_n(\mathbf{X}_{n+j}), \mathbf{Y}_{n+j}) - l(\widehat{g}_n(\mathbf{Z}_{n+j}), \mathbf{Y}_{n+j})$

Decomposition

- T_2 converges to zero in probability for **good estimators** (peak performance for black-box models)

$$\begin{aligned} T_2 &= R(\hat{f}_n) - R(f^*) + R_S(g^*) - R_S(\hat{g}_n) \\ &\leq \max\{R(\hat{f}_n) - R(f^*), R_S(\hat{g}_n) - R_S(g^*)\} \quad \underbrace{= O_P(n^{-\gamma})}_{\text{reasonable assumption}} \end{aligned}$$

In the literature, the convergence rate $\gamma > 0$ has been extensively investigated [Wasserman, 2006, Schmidt-Hieber et al., 2020].

- T_3 is related to H_0

$$T_3 = R(f^*) - R(g^*) = 0, \quad \text{under } H_0$$

Motivation

- **Main idea:**

- **Normalize** T_1 by its standard derivation, which can be estimated by a sample standard derivation of evaluations on the inference set. Then, the normalized T_1 follows $N(0, 1)$ asymptotically by CLT.
- After normalization, T_2 is **convergence in probability** when $n \rightarrow \infty$, and $T_3 = 0$ under H_0 .

- Consider the following test statistic:

$$\frac{\sqrt{m}}{\hat{\sigma}_n} (\hat{R}(\hat{f}_n) - \hat{R}_S(\hat{g}_n)) = \frac{\sum_{j=1}^m \Delta_{n,j}}{\sqrt{m}\hat{\sigma}_n} = \frac{\sqrt{m}}{\hat{\sigma}_n} T_1 + \frac{\sqrt{m}}{\hat{\sigma}_n} T_2 + \frac{\sqrt{m}}{\hat{\sigma}_n} T_3,$$

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- Asymptotically Normally Distributed? **It may be WRONG!!**

Bias-sd-ratio issue

One unusual issue for the test statistic is **vanishing** standard deviation:

Under H_0 , if $\hat{f}_n \xrightarrow{P} f^*$, $\hat{g}_n \xrightarrow{P} g^*$, and $f^* = g^*$, then $\hat{\sigma}_n \xrightarrow{P} 0$

Issues:

- **CLT may not hold for T_1 .** CLT requires a standard deviation is fixed, or bounded away from zero.
- **Bias-sd-ratio.** Both bias and sd are convergence to zeros:

$$\frac{\sqrt{m}T_2}{\hat{\sigma}_n} = \sqrt{m} \left(\frac{R(\hat{f}_n) - R(f^*) + R_S(g^*) - R_S(\hat{g}_n)}{\hat{\sigma}_n} \right) = \sqrt{m} \left(\frac{\text{bias} \xrightarrow{P} 0}{\text{sd} \xrightarrow{P} 0} \right).$$

- If T_2 and $\hat{\sigma}_n$ are in the same order, $\sqrt{m}\hat{\sigma}_n^{-1}T_2 = O_P(\sqrt{m})$, **kills the null distribution.**

Solution

- The issue is caused by **vanishing** standard deviation, we can address it by **perturbation**.
- **One-split test**. The proposed test statistic is given as:

$$\Lambda_n^{(1)} = \frac{\sum_{j=1}^m \Delta_{n,j}^{(1)}}{\sqrt{m\hat{\sigma}_n}}, \quad \Delta_{n,j}^{(1)} = \Delta_{n,j} + \rho_n \varepsilon_j, \quad (3)$$

where $\hat{\sigma}_n$ is the sample standard derivation based on $\{\Delta_{n,j}^{(1)}\}_{j=1}^m$ conditional on \hat{f}_n and \hat{g}_n , $\rho_n \rightarrow \rho$ is a level of perturbation.

- Note that $\hat{\sigma}_n^{(1)} \rightarrow \sigma^{(1)} \geq \rho > 0$.

Decomposition

Reconsider the decomposition of $\Lambda_n^{(1)}$:

$$\Lambda_n^{(1)} = \underbrace{\frac{\sqrt{m}}{\hat{\sigma}_n^{(1)}} \left(\frac{1}{m} \sum_{j=1}^m (\Delta_{n,j}^{(1)} - \mathbb{E}(\Delta_{n,j}^{(1)} | \mathcal{E}_n)) \right)}_{\rightarrow N(0,1) \text{ by conditional CLT of triangular array}}$$
$$+ \underbrace{\frac{\sqrt{m}}{\hat{\sigma}_n^{(1)}} \left(R(\hat{f}_n) - R(f^*) - (R_S(\hat{g}_n) - R_S(g^*)) \right)}_{= O_p(m^{1/2}n^{-\gamma}) \text{ by prediction consistency}} + \underbrace{\frac{\sqrt{m}}{\hat{\sigma}_n^{(1)}} (R(f^*) - R_S(g^*))}_{= 0 \text{ under } H_0}.$$

- If the **splitting condition** $m^{1/2}n^{-\gamma} = o_p(1)$ is satisfied, then

$$\Lambda_n^{(1)} \xrightarrow{d} N(0, 1) \text{ under } H_0$$

Asymptotic null distribution

- **Assumption A** (Prediction consistency). For some constant $\gamma > 0$, (\hat{f}_n, \hat{g}_n) satisfies

$$(R(\hat{f}_n) - R(f^*)) - (R_S(\hat{g}_n) - R_S(g^*)) = O_p(n^{-\gamma}). \quad (4)$$

- **Assumptions B-C** are standard assumptions for CLT under triangle arrays [Cappé et al., 2006].

Theorem 2 (Asymptotic null distribution of $\Lambda_n^{(1)}$)

In addition to Assumptions A, B, and C, if $m = o(n^{2\gamma})$, then under H_0 ,

$$\Lambda_n^{(1)} \xrightarrow{d} N(0, 1), \quad \text{as } n \rightarrow \infty. \quad (5)$$

According to the asymptotic null distribution of $\Lambda_n^{(1)}$ in Theorem 2, we calculate the p -value $P^{(1)} = \Phi(\Lambda_n^{(1)})$.

Power analysis

Consider an alternative hypothesis $H_a : R(f^*) - R_S(g^*) = -m^{-1/2}\delta < 0$ for $\delta > 0$. The power functions of the one-split test and its combined test can be written as

$$\pi_n(\delta) = \mathbb{P}(P^{(1)} \leq \alpha | H_a), \quad \bar{\pi}_n(\delta) = \mathbb{P}(\bar{P}^{(1)} \leq \alpha | H_a).$$

Theorem 3 (Local limiting power of the one-split test)

Suppose that the one-split test (3) satisfies Assumptions A-C and $m = o(n^{2\gamma})$, then

$$\liminf_{n \rightarrow \infty} \pi_n(\delta) = \Phi\left(\frac{\delta}{\sigma(1)} - z_\alpha\right), \quad \text{and} \quad \lim_{\delta \rightarrow \infty} \liminf_{n \rightarrow \infty} \pi_n(\delta) = 1, \quad (6)$$

where $z_\alpha = \Phi^{-1}(1 - \alpha)$ is the z-multiplier of the standard normal distribution.

Splitting condition

- **Question:** How to determine the estimation / inference ratio?
 $m = o(n^{2\gamma})$ for an **unknown** $\gamma > 0$.
- **Log-ratio sample splitting scheme.** Specifically, given a sample size $N \geq N_0$, the estimation and inference sizes n and m are obtained:

$$n = \lceil x_0 \rceil, \quad m = N - n,$$

$$\text{where } x_0 \text{ is a solution of } \left\{ x + \frac{N_0}{2 \log(N_0/2)} \log(x) = N \right\}. \quad (7)$$

- **Splitting ratio condition** is automatically satisfied!

Lemma 4 (Log-ratio sample splitting scheme)

The estimation and inference sample sizes (n, m) , determined by the log-ratio sample splitting formula (7), satisfies $m = o(n^{2\gamma})$ for any $\gamma > 0$ in Assumption A.

More comments

- **Power.** Heuristic data-adaptive sample splitting scheme.
- **Two-split test.** One-split test is valid for any perturbation $\rho > 0$, if you don't like a custom parameter, use **two-split test** (further splitting an inference sample into two equal subsamples yet the perturbation is not required).
- **CV.** Combining p-values over repeated random splitting.

Algorithm

Algorithm 1 One-split test for feature relevance to prediction

- 1: **Input:** Data: $(\mathbf{x}_i, \mathbf{y}_i)_{i=1}^N$; Set of hypothesized feats: \mathcal{S} ; #splitting: U
 - 2: **Output:** p -value for testing (2)
 - 3: Determine the splitting ratio ξ and the perturbation level ρ (log-ratio or data-adaptive scheme)
 - 4: **for** $u = 1, \dots, U$ **do**
 - 5: Shuffle the data
 - 6: Split the data into estimation / inference sets, where $m = \hat{\zeta}N$ and $n = N - m$
 - 7: Compute $\Lambda_u^{(1)}$ from (3)
 - 8: Compute p -value $P_u^{(1)} = \Phi(\Lambda_u^{(1)})$
 - 9: **end for**
 - 10: Compute the combined p -value $\bar{P}^{(1)}$
-

- Just fit a DL model U -times, U can be as small as 1.

Numerical experiments

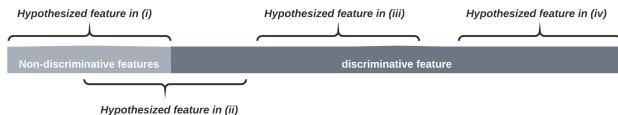
- **Comparison with existing black-box tests.**

- Sample size $N = 1000$, dimension $p = 5$.
- $\mathbf{X} = (X_1, \dots, X_5)^\top$ follows a uniform distribution on $[-1, 1]$ with a pairwise correlation $\rho_{ij} = 0.5^{|i-j|}$.
- $Y = 0.02(X_1 + X_2 + X_3) + 0.05\epsilon$

Test	Return	H_0	
One-split	p -value	risk-invariance $R(f^*) = R_S(g^*)$,	0.003
Two-split	p -value	risk-invariance $R(f^*) = R_S(g^*)$	0.018
HRT	p -values for all feats	conditional indep $\mathbf{X}_j \perp \mathbf{Y} \mathbf{X}_{-j}$	(0.840, 0.045, 0.064, 0.900, 0.158)
LOCO	p -values for all feats	equal errors with/without feat j for a given dataset	(0.132, 0.791, 0.180, 0.435, 0.342)
PT	p -value	marginal indep $\mathbf{X}_S \perp \mathbf{Y}$	0.010
HPT	p -value	marginal indep $\mathbf{X}_S \perp \mathbf{Y}$	0.001

Numerical experiments

- Simulation for a neural network: $Y = f^*(\mathbf{X}) + \epsilon$.
 - $f^*(\mathbf{x})$ is a neural network. $\mathbf{X} \sim N(\mathbf{0}, B\Sigma)$, $\Sigma_{ij} = r^{|i-j|}$, $r \in [0, 1)$
 - $f^*(\mathbf{x}) = g^*(\mathbf{z})$ only depends on a subset of features of \mathbf{x} , in which $\mathbf{z}_{S_0} = \mathbf{0}$ and $\mathbf{z}_{S_0^c} = \mathbf{x}_{S_0^c}$ with $S_0 = \{1, \dots, |S_0|\}$.
 - Given a hypothesized index set \mathcal{S} , our goal is to test if $\mathbf{X}_{\mathcal{S}}$ is relevant to predicting the outcome Y .



- (i) $\mathcal{S} \cup S_0 = S_0$ for Type I error. (ii)-(iv): $\mathcal{S} \cup S_0 \neq S_0$ for power.
- (ii) \rightarrow (iv), the distance (or correlation) between \mathcal{S} and S_0 is increasing (or decreasing), thus the power is expected to go up.

Numerical experiments

Example 1. (*Impact of the sample size*) This example (Table 1) concerns the performance of the proposed tests in relation to the sample size N based on *data-adaptive* tuning methods, where N ranges from 2000 to 10000, $B = 0.4$, $r = 0.25$, $p = 100$, $\varpi = 128$, $\tau = 2$, $L = 3$, $|\mathcal{S}_0| = 5$.

Test	Sample size	Type I error	Power	Time (Second)
One-split	2000	0.043	(0.25, 0.79, 0.85)	15.2(0.1)
	6000	0.050	(0.61, 0.99, 1.00)	41.2(0.3)
	10000	0.049	(0.89, 1.00, 1.00)	66.0(0.4)
Two-split	2000	0.050	(0.11, 0.26, 0.31)	14.0(0.1)
	6000	0.035	(0.18, 0.51, 0.58)	37.0(0.2)
	10000	0.040	(0.19, 0.77, 0.75)	61.6(0.4)
Comb. one-split	2000	0.034	(0.26, 1.00, 0.95)	37.9(0.1)
	6000	0.046	(0.86, 1.00, 1.00)	68.3(0.3)
	10000	0.045	(1.00, 1.00, 1.00)	107.2(0.7)
Comb. two-split	2000	0.015	(0.09, 0.26, 0.29)	38.0(0.1)
	6000	0.030	(0.10, 0.70, 0.65)	76.3(0.5)
	10000	0.014	(0.13, 0.93, 0.92)	110.3(0.5)

Table 1: Empirical Type I errors and powers of the one-split and two-split tests, their combined tests in Example 1 at a nominal level $\alpha = 0.05$.

Numerical experiments

Example 2. (*Impact of the strength of features of interest*) This example (Table 2) concerns the performance of the proposed tests with respect to the magnitude of hypothesized features B , where $B = 0.2, 0.4, 0.6$, $N = 6000$, $p = 100$, $r = 0.25$, $\varpi = 128$, $\tau = 2$, $L = 3$, and $|\mathcal{S}_0| = 5$.

Test	B	Type I error	Power
One-split	0.2	0.057	(0.24, 0.68, 0.78)
	0.4	0.050	(0.61, 0.99, 1.00)
	0.6	0.057	(0.97, 1.00, 1.00)
Two-split	0.2	0.049	(0.06, 0.12, 0.14)
	0.4	0.035	(0.18, 0.51, 0.58)
	0.6	0.041	(0.37, 0.97, 0.98)
Comb. one-split	0.2	0.027	(0.27, 0.93, 0.93)
	0.4	0.046	(0.86, 1.00, 1.00)
	0.6	0.033	(1.00, 1.00, 1.00)
Comb. two-split	0.2	0.019	(0.00, 0.00, 0.03)
	0.4	0.030	(0.10, 0.70, 0.65)
	0.6	0.012	(0.45, 1.00, 1.00)

Table 2: Empirical Type I errors and powers of the one-split and two-split tests, and their combined tests in Example 2 at a nominal level $\alpha = 0.05$. The data-adaptive tuning scheme is applied.

Numerical experiments

- **Example 3.** (*Impact of the depth and width of a neural network*) This example concerns the performance of the proposed tests in terms of the width ϖ and depth L of a neural network, where $N = 6000$, $L = 2, 3, 4$, $\varpi = 32, 64, 128$, $B = 0.4$, $r = 0.25$, $p = 100$, $\tau = 2$, $L = 3$, and $|\mathcal{S}_0| = 5$.
- **Example 4.** (*Impact of the number of hypothesized features*) This example concerns the proposed tests with respect to the number of hypothesized features $|\mathcal{S}_0| = 3, 5, 10$.
- **Example 5.** (*Impact of feature correlations*) This example concerns the proposed tests in terms of the feature correlation $r = .2, .4, .6$.
- **Example 6.** (*Impact of different modes of combining p-values*) This example concerns the combined tests with different ways of combining p-values, including the Hommel, the Bonferroni, the first quantile, the median, the Cauchy, and the harmonic methods.

Numerical experiments

• Role of perturbation.

- $\mathcal{S}_0 = \{1, 2, 3\}$, $\mathbf{X} \sim N(\mathbf{0}, B\Sigma)$, $\Sigma_{ij} = r^{|i-j|}$, $r \in [0, 1)$
- $\Sigma_{1j} = \Sigma_{j1} = .1$; $j = 1, \dots, p$, and $\Sigma_{ij} = 0$, if $i, j \neq 1$ and $i \neq j$.
- Only partial features are observed in a sample $(\mathbf{x}_i^{(N)}, y_i^{(N)})_{i=1}^N$, where $\mathbf{x}_i^{(N)} = (\mathbf{x}_{i1}, \dots, \mathbf{x}_{id_N})^\top$ and $y_i^{(N)} = f^*(\tilde{\mathbf{x}}_i^{(N)}) + \epsilon_i$, $d_N \rightarrow d$ as $N \rightarrow \infty$, and $\tilde{\mathbf{x}}_i^{(N)} = (\mathbf{x}_{i1}, \dots, \mathbf{x}_{id_N}, 0, \dots, 0)^\top$ is a d -dimensional vector.

Test	$N = 2000$	$N = 6000$	$N = 10000$
One-split without perturbation	0.083	0.109	0.193
One-split with perturbation	0.057	0.053	0.061
Two-split	0.048	0.051	0.047

Table 3: Type I errors of the one-split tests with and without perturbation and the two-split test in Section 6.4 at a nominal level $\alpha = 0.05$.

Summary

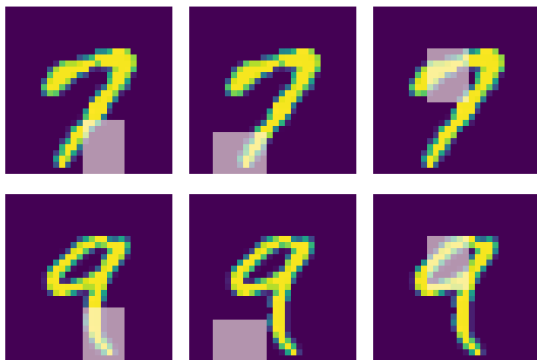
- Summarize the simulation results.

		<i>Advantage</i>	Evidence
Test	One-split	<i>More powerful</i>	Tables 3-5
	Two-split	<i>No need to perturb data</i>	Equation (14)
Combine	Comb.	<i>More powerful</i>	Tables 3-5
	Non-comb.	<i>Less computation time</i>	Table 3
Ratio	Data-adaptive	<i>More powerful</i>	Tables 3-5
	Log-ratio	<i>No need to tune the ratio, and less computation time</i>	Lemma 4, Table 3

Table 4: Advantage for different tests, combining, and tuning methods.

Real application

- The MNIST handwritten digits dataset [LeCun et al., 1998]. In particular, we extract 14,251 images from the dataset with labels '7' and '9' to discriminate between these two digits.



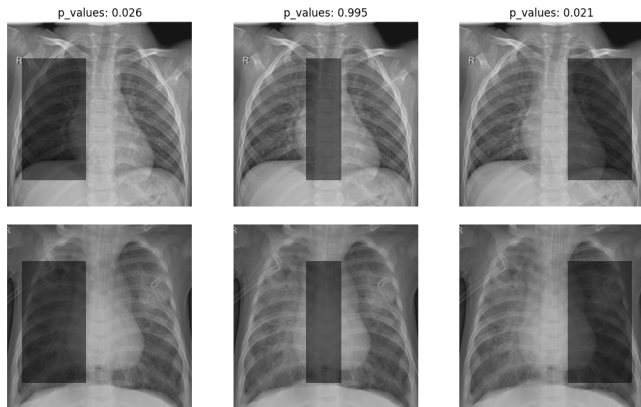
Real application

Test	p -values (case 1, case 2, case 3)	Time(Second)
One-split	(0.174, 0.329, 0.000)	4289
Two-split	(0.959, 0.569, 0.000)	4772
Comb. one-split	(0.385, 1.000, 0.000)	11404
Comb. two-split	(0.544, 0.192, 0.000)	13060

Table 5: P-values and runtimes of the one-split and two-split tests, their combined tests, and the permutation test in the MNIST benchmark example at a nominal level $\alpha = 0.05$.

Real application

- pneumonia diagnosis dataset [Keremany et al., 2018]. This dataset consists of 5,863 X-ray images, each labeled as “Pneumonia” or “Normal.”



Real application

Test	p -values (case 1, case 2, case 3)	Time(Second)
One-split	(0.026, 0.995, 0.021)	15242
Two-split	(0.212, 0.561, 0.065)	14020
Comb. one-split	(0.041, 0.635, 0.075)	64416
Comb. two-split	(0.053, 0.754, 0.084)	64761


Table 6: P-values and runtimes of the one-split and two-split tests, and their combined tests in the chest X-ray dataset at a nominal level $\alpha = 0.05$.

Contribution




- A **novel risk-based hypothesis** is proposed in (2), as well as its relation to conditional independence tests.
- We derive the **one-split/two-split tests** based on the differenced empirical loss with and without hypothesized features. Theoretically, we show that the one-split and two-split tests, as well as their combined tests, can **control the Type I error** while being **consistent in terms of power**;
- The proposed tests only require a **limited number of refitting**, and we develop the Python library `dnn-inference` and examine the utility of the proposed tests on various simulated examples and two real datasets.

Thank you!




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