

MOTIVATION

Biometric identification = checks correspondance of two measurements (x, x') .

Given a similarity s and a threshold t ,

$$(x, x') \text{ is a match} \Leftrightarrow s(x, x') > t. \quad (1)$$

The ROC curve of s gives the true positive rate (TPR) given the false positive rate (FPR) associated to eq. (1) for all thresholds $t \in \mathbb{R}$.

An usual approach is to optimize the Area under the ROC_s curve (AUC) of the similarity function s .

Biometric systems are deployed for a fixed, low FPR, which is hard to optimize in practice (see [1]).

CONTRIBUTIONS

An extension of the algorithm TreeRank (see [2]) that learns a symmetric similarity function which optimizes the ROC curve for the similarity ranking problem (see [1]).

Statistical guarantees in $\|\cdot\|_\infty$ in the ROC space.

Empirical illustration on synthetic data.

Trials on real data.

PRELIMINARIES

Classification setting. Assume $(X, Y) \sim P$, with:

- $Y \in \{1, \dots, K\}$ the output label,
- $X \in \mathcal{X} \subset \mathbb{R}^d$ input random variable.

Similarity learning. Select similarity function S s.t.

the larger $S(X, X')$ the higher $\mathbb{P}\{Y = Y' \mid X, X'\}$,

with $(X, Y) \perp (X', Y') \sim P$.

Optimal similarities \mathcal{S}^* are increasing transforms of:

$$\eta(x, x') = \mathbb{P}\{Y = Y' \mid (X, X') = (x, x')\}.$$

The accuracy of any $s \in \mathcal{S}$ can be measured by:

$$d_p(s, s^*) = \|\text{ROC}_s - \text{ROC}^*\|_p,$$

where $s^* \in \mathcal{S}^*$ and $p \in [1, +\infty]$.

Given i.i.d. copies $\mathcal{D}_n = \{(X_i, Y_i)\}_{i=1}^n$ of (X, Y) , one wants to rank the dependent pairs

$$\{(X_i, X_j), Z_{i,j}\} : 1 \leq i < j \leq n\},$$

TreeRank (see [2].) Recursive technique that builds a piecewise constant score s_{D_n} for bipartite ranking.

Bipartite ranking considers data $\{(X_i, Y_i)\}_{i=1}^n$, i.i.d. copies of $(X, Y) \in \mathcal{X} \times \{-1, +1\}$.

TreeRank splits the input space recursively for weighted classification with a class $\mathcal{A} \subset \mathcal{P}(\mathcal{X})$.

Under assumptions on the distribution and \mathcal{A} , [2] (Corollary 16 therein), state that, given a tree of depth $D = D_n \sim \log(\sqrt{n})$ as $n \rightarrow \infty$, $\forall \delta > 0$, $\exists \lambda$ s. t., w.p. $\geq 1 - \delta$, $\forall n \in \mathbb{N}$, for $i \in \{1, +\infty\}$,

$$d_i(s_{D_n}, s^*) \leq \exp(-\lambda \sqrt{\log n}). \quad (2)$$

SIMILARITY TREERANK

For $F_\sigma(\mathcal{C})$, $\sigma \in \{-, +\}$ and $\mathcal{C} \subset \mathcal{X} \times \mathcal{X}$, introduce :

$$\widehat{F}_{\sigma,n}(\mathcal{C}) = \frac{1}{n_\sigma} \sum_{i < j} \mathbb{I}\{(X_i, X_j) \in \mathcal{C}, Z_{i,j} = \sigma\},$$

with $n_\sigma = (2/(n(n-1))) \sum_{i < j} \mathbb{I}\{Z_{i,j} = \sigma\}$, which are not i.i.d. averages, but ratios of U -statistics of degree two (i.e. averages over pairs of observations, see [1]).

Input. Maximal depth $D \geq 1$, class \mathcal{A} of measurable symmetric subsets of $\mathcal{X} \times \mathcal{X}$, training dataset \mathcal{D}_n .

1. (INITIALIZATION.) Set $\mathcal{C}_{0,0} = \mathcal{X} \times \mathcal{X}$, $\alpha_{d,0} = \beta_{d,0} = 0$ and $\alpha_{d,2^d} = \beta_{d,2^d} = 1$ for $d \geq 0$.

2. (ITERATIONS.)

For $d = 0, \dots, D-1$ and $k = 0, \dots, 2^d - 1$:

a) (OPTIMIZATION STEP.)

Set the entropic measure:

$$\Lambda_{d,k+1}(\mathcal{C}) = (\alpha_{d,k+1} - \alpha_{d,k}) \widehat{F}_{+,n}(\mathcal{C}) - (\beta_{d,k+1} - \beta_{d,k}) \widehat{F}_{-,n}(\mathcal{C}).$$

Find the best subset $\mathcal{C}_{d+1,2k}$ of the cell $\mathcal{C}_{d,k}$ in the AUC sense:

$$\mathcal{C}_{d+1,2k} = \operatorname{argmax}_{\mathcal{C} \in \mathcal{A}, \mathcal{C} \subset \mathcal{C}_{d,k}} \widehat{\Lambda}_{d,k+1}(\mathcal{C}).$$

Then, set $\mathcal{C}_{d+1,2k+1} = \mathcal{C}_{d,k} \setminus \mathcal{C}_{d+1,2k}$.

b) (UPDATE.) Set

$$\alpha_{d+1,2k+1} = \alpha_{d,k} + \widehat{F}_{-,n}(\mathcal{C}_{d+1,2k}),$$

$$\beta_{d+1,2k+1} = \beta_{d,k} + \widehat{F}_{+,n}(\mathcal{C}_{d+1,2k}),$$

and $\alpha_{d+1,2k+2} = \alpha_{d,k+1}$, $\beta_{d+1,2k+2} = \beta_{d,k+1}$.

3. (OUTPUT.) After D iterations, get the piecewise constant similarity function:

$$s_D(x, x') = \sum_{k=0}^{2^D-1} (2^D - k) \mathbb{I}\{(x, x') \in \mathcal{C}_{D,k}\}.$$

GUARANTEES

The theoretical guarantees formulated in eq. (2) for the standard bipartite ranking setup remain valid in the similarity learning framework.

The following technical assumptions are involved:

- the feature space \mathcal{X} is bounded;
- $\alpha \mapsto \text{ROC}^*(\alpha)$ is twice differentiable with a bounded first order derivative;
- the class \mathcal{A} is intersection stable, i.e. $\forall (\mathcal{C}, \mathcal{C}') \in \mathcal{A}^2, \mathcal{C} \cap \mathcal{C}' \in \mathcal{A}$;
- the class \mathcal{A} has finite VC dimension $V < +\infty$;
- we have $\{(x, x') \in \mathcal{X}^2 : \eta(x, x') \geq q\} \in \mathcal{A}$ for any $q \in [0, 1]$;

Let s_D denote the output of the SIMILARITY TREERANK algorithm with depth D .

Theorem 1. Assume that the conditions above are fulfilled. Choose $D = D_n$ so that $D_n \sim \sqrt{\log n}$, as $n \rightarrow \infty$. Then, for all $\delta > 0$, there exists a constant λ s.t., with probability at least $1 - \delta$, we have for all $n \geq 2$:

$$d_\infty(s_{D_n}, s^*) \leq \exp(-\lambda \sqrt{\log n}).$$

SYNTHETIC EXPERIMENTS

We generate data with a random tree of depth D_{gt} and fix $\mathcal{X} = \mathbb{R}^3$, $\delta = 0.01$, $n_{\text{test}} = 100,000$, $n_{\text{train}} = 150 \cdot (5/4)^{D_{gt}}$. TreeRank outputs s_D , results feature 95% CI's based on 400 runs.

Class asymmetry		
p_+	$D_1(s_D, s^*)$	$D_\infty(s_D, s^*)$
0.5	0.07(±0.07)	0.30(±0.07)
10^{-1}	0.08(±0.08)	0.31(±0.08)
10^{-3}	0.42(±0.17)	0.75(±0.17)
$2 \cdot 10^{-4}$	0.45(±0.08)	0.81(±0.08)

Parameters: $D = D_{gt} = 3$.

Model bias		
D	$D_1(s_D, s^*)$	$D_\infty(s_D, s^*)$
1	0.21(±0.13)	0.65(±0.13)
2	0.11(±0.10)	0.43(±0.10)
3	0.07(±0.07)	0.30(±0.07)
8	0.06(±0.06)	0.28(±0.06)

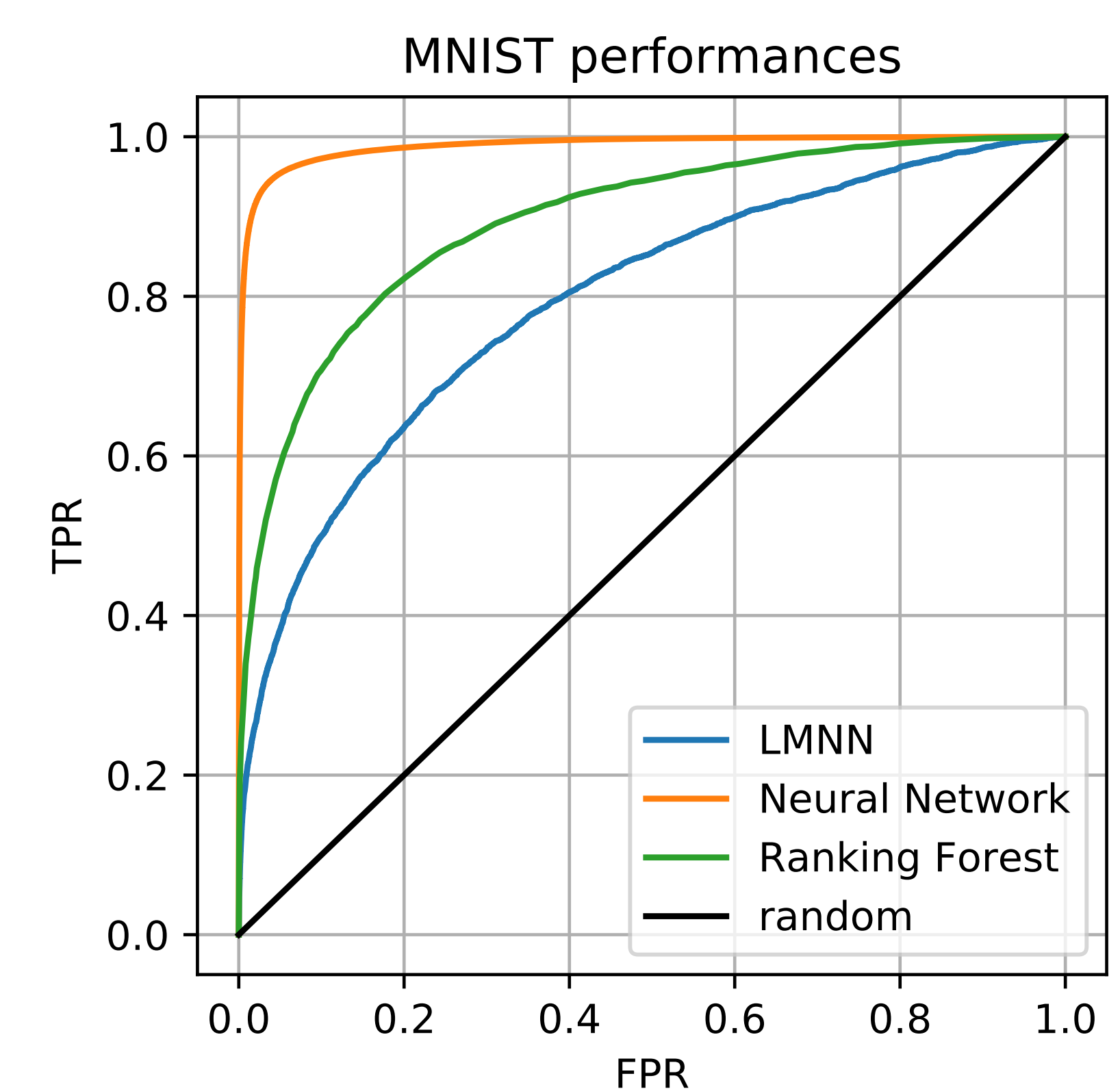
Parameters: $D_{gt} = 3, p = 0.5$.

This table illustrate two parameters that impair generalization: pronounced **class asymmetry** and **under-specified models**.

REAL DATA EXPERIMENTS

We try to validate our model by learning a similarity on MNIST, reduced by PCA. We compare three models:

- A linear metric learning algorithm: LMNN [3],
- A simple metric on a neural network encoding, trained for classification,
- Similarity TreeRank with decision stumps as \mathcal{A} .



Conclusions: Limited competitiveness of TreeRank. \mathcal{A} is not expressive enough ?

Future work will explore Treerank approaches with neural networks for \mathcal{A} .

REFERENCES

- [1] Robin Vogel, Aurélien Bellet, and Stéphan Cléménçon. A probabilistic theory of supervised similarity learning for pointwise ROC curve optimization. In *ICML*, 2018.
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- [3] Kilian Q. Weinberger and Lawrence K. Saul. Distance Metric Learning for Large Margin Nearest Neighbor Classification. *JMLR*, 10:207–244, 2009.