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# Efficient Direct Density Ratio Estimation for Non-stationarity Adaptation and Outlier Detection

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# Density Ratio

$$\left. \begin{array}{l} x_1, \dots, x_n \sim_{i.i.d.} p(x) \\ x'_1, \dots, x'_m \sim_{i.i.d.} q(x') \end{array} \right\} \xRightarrow{\text{estimate}} w(x) = \frac{p(x)}{q(x)}, \quad \text{Density Ratio (Importance)}$$

Density ratio can be used for various succeeding tasks:

Feature selection (Suzuki, et al., ECML workshop 2008),

Multi-task learning (Bickel, et al., ICML 2008),

Domain Adaptation (Storkey and Sugiyama, NIPS 2006, Tsuboi et al., SDM 2008), etc.

Our main applications: **Covariate shift adaptation, Outlier detection**

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# Estimation of Density Ratio

**Samples :**

$$x_1^{(\text{tr})}, \dots, x_n^{(\text{tr})} \sim p^{(\text{tr})}(x),$$
$$x_1^{(\text{te})}, \dots, x_m^{(\text{te})} \sim p^{(\text{te})}(x)$$

- **dim  $x$  is small  $\implies$  Naive methods are available, e.g. separately estimate  $\hat{p}^{(\text{tr})}, \hat{p}^{(\text{te})}$  by kernel density estimator, and obtain  $\hat{w}(x) = \hat{p}^{(\text{te})} / \hat{p}^{(\text{tr})}$ .**
- **dim  $x$  is not small  $\implies$  direct density ratio estimation**
  - **Kernel Mean Matching** (Huang, et al., NIPS2006): mean matching in RKHS.
  - **Logistic Regression** (Bickel et al., ICML2008): binary classification approach
  - **KLEIP** (Sugiyama, et al., NIPS2007): inference using KL-divergence
  - **Proposed Method: Least squares approach**
    - \* **Efficient computation of estimator and leave-one-out cross validation**

# Least Squares Approach to Importance Estimation

Square error of density ratio:

$$\begin{aligned} & \frac{1}{2} \int \left( w(x) - \frac{p^{(\text{te})}(x)}{p^{(\text{tr})}(x)} \right)^2 p^{(\text{tr})}(x) dx \\ &= \frac{1}{2} \int w(x)^2 p^{(\text{tr})}(x) dx - \int w(x) p^{(\text{te})}(x) dx + (\mathbf{const.}) \end{aligned}$$

note: In succeeding tasks,  $w(x^{(\text{tr})})$  is often used. Thus, the expectation with  $p^{(\text{tr})}$  is valid.

**Estimator:**  $w(x) = \sum_{\ell=1}^b \alpha_{\ell} \varphi_{\ell}(x)$ ,  $\varphi_{\ell}(x) = e^{-\gamma \|x - c_{\ell}\|^2} > 0$ ,  $c_{\ell}$ : kernel center

**empirical loss:**  $\frac{1}{2n} \sum_{i=1}^n w(x_i^{(\text{tr})})^2 - \frac{1}{m} \sum_{j=1}^m w(x_j^{(\text{te})}) = \frac{1}{2} \alpha^{\top} H \alpha - g^{\top} \alpha \longrightarrow \min_{\alpha}$

$$H_{\ell\ell'} := \frac{1}{n} \sum_{i=1}^n \varphi_{\ell}(x_i^{(\text{tr})}) \varphi_{\ell'}(x_i^{(\text{tr})}), \quad g_{\ell} := \frac{1}{m} \sum_{j=1}^m \varphi_{\ell}(x_j^{(\text{te})})$$

# Non-Negativity Condition and Regularization

Impose the constraint  $w(x) \geq 0$  to the estimator:

$$(I) \quad \min_{\alpha} \frac{1}{2} \alpha^{\top} H \alpha - g^{\top} \alpha + \lambda R(\alpha) \rightarrow \tilde{\alpha}, \quad \hat{\alpha}_{\ell} = \max\{\tilde{\alpha}_{\ell}, 0\}$$

$$(II) \quad \min_{\alpha} \frac{1}{2} \alpha^{\top} H \alpha - g^{\top} \alpha + \lambda R(\alpha), \text{ s.t. } \alpha \geq 0 \rightarrow \hat{\alpha}$$

■ **Proposed estimator: (I) with  $L_2$ -regularization  $R(\alpha) = \|\alpha\|_2^2$  called **uLSIF** (unconstrained Least-square Importance Inference)**

• **Estimator  $\hat{\alpha}$  is analytically computed** :  $\hat{\alpha} = \max\{(H + \lambda I)^{-1}g, 0\}$

• **Leave-one-out cross validation is analytically computed.**

$$\text{LOOCV} = \frac{1}{n} \sum_{i=1}^n \left[ \frac{1}{2} (\hat{w}^{(i)}(x_i^{(\text{tr})}))^2 - \hat{w}^{(i)}(x_i^{(\text{te})}) \right] : \text{directly computed via } \hat{\alpha}.$$

$\hat{w}^{(i)}$  : estimator obtained without the samples  $x_i^{(\text{tr})}, x_i^{(\text{te})}$  ( $i = 1, \dots, n, n \leq m$ ).

■ **(II) +  $L_1$ -regularization  $\implies$  regularization path in term of  $\lambda$  is obtained. Numerically rather unstable.**

# Relation between Proposed and Existing Methods

Methods	Density estimation	Optimization	Out-of-sample prediction (CV)
KDE	Necessary	<b>Analytic</b>	<b>Possible</b>
KMM	<b>Not necessary</b>	Convex quadratic	Not possible
LogReg	<b>Not necessary</b>	Convex non-linear	<b>Possible</b>
KLIEP	<b>Not necessary</b>	Convex non-linear	<b>Possible</b>
uLSIF	<b>Not necessary</b>	<b>Analytic</b>	<b>Possible</b>

- KMM estimates not the function  $w(x)$  but the values  $w(x_i^{(\text{tr})})$ ,  $i = 1, \dots, n$ . Thus, cross validation will not be possible.
- Model selection of uLSIF is computationally much more efficient than LogReg and KLIEP, because analytic formula of leave-one-out cv is available.

# Numerical Examples: Estimation of Density Ratio

- **Estimate**  $w(x) = p^{(\text{te})}(x)/p^{(\text{tr})}(x)$ .

$$x_1^{(\text{tr})}, \dots, x_n^{(\text{tr})} \sim N_d(\mathbf{0}, \mathbf{I}_d)$$

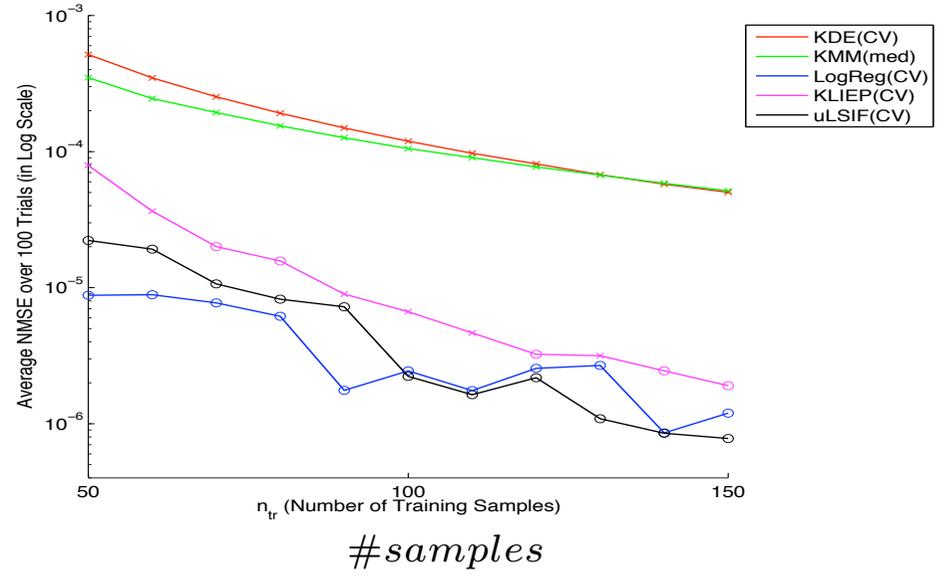
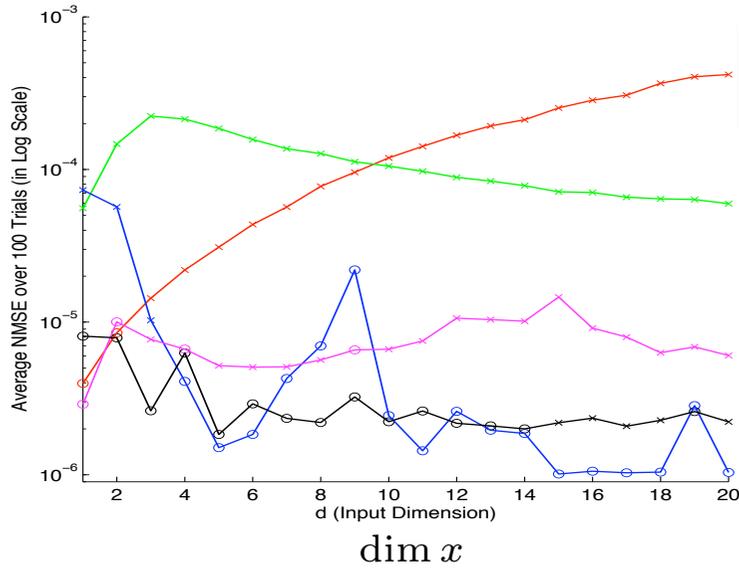
$$x_1^{(\text{te})}, \dots, x_{1000}^{(\text{te})} \sim N_d(\mathbf{e}_1, \mathbf{I}_d), \quad \mathbf{e}_1 = (1, 0, \dots, 0)^\top \in \mathbb{R}^d$$

- **setup 1:**  $d = 1, 2, \dots, 20, \quad n = 100$
- **setup 2:**  $n = 50, 60, \dots, 150, \quad d = 10$

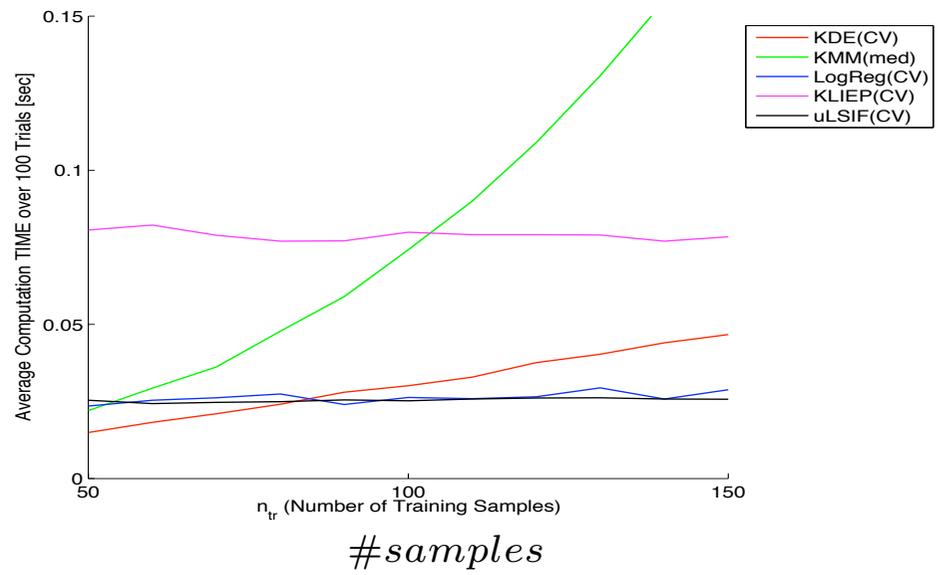
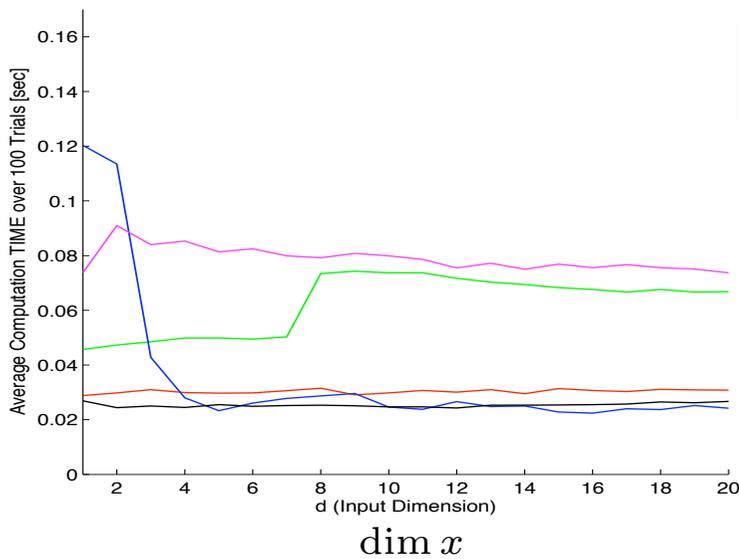
- **Evaluation of accuracy:** square error on  $x_1^{(\text{tr})}, \dots, x_n^{(\text{tr})}$

$$\text{normalized-MSE (NMSE)} = \frac{1}{n} \sum_{i=1}^n \left( \frac{\hat{w}(x_i^{(\text{tr})})}{\sum_{j=1}^n \hat{w}(x_j^{(\text{tr})})} - \frac{w(x_i^{(\text{tr})})}{\sum_{j=1}^n w(x_j^{(\text{tr})})} \right)^2$$

# Square Errors of Density Ration Estimation

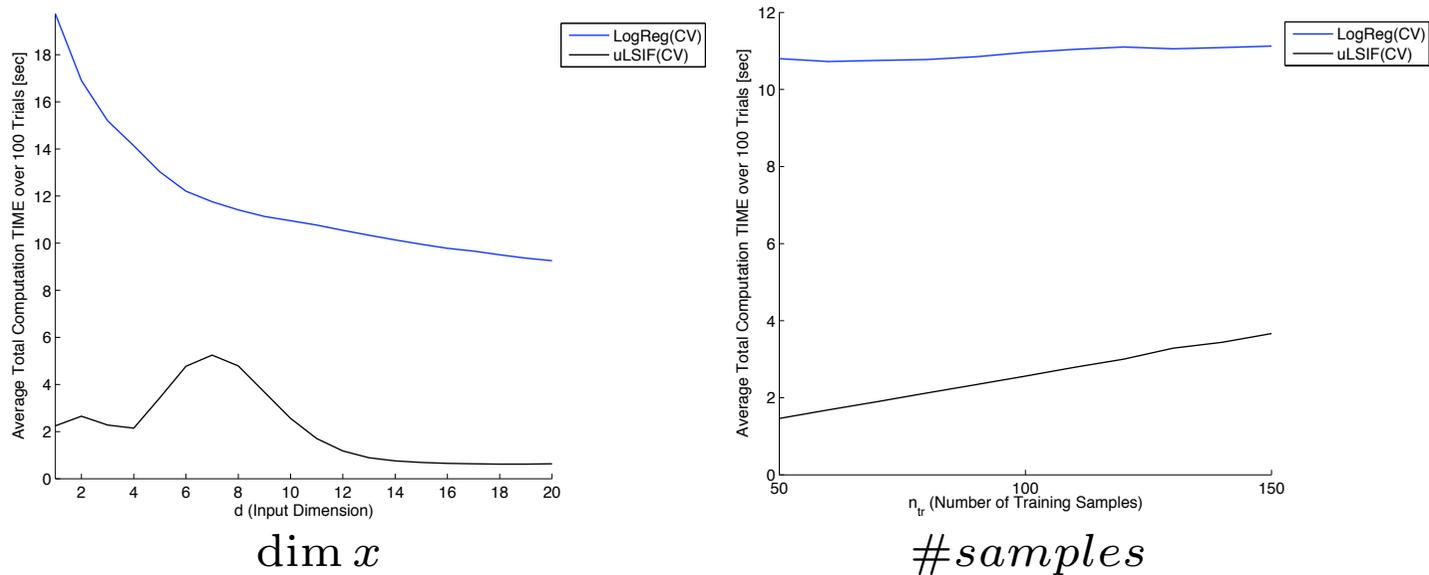


# Computation Time for Fixed Model Parameters



- Least-square error: uLSIF, Logistic reg., KLIEP < KDE, KMM
- Computation time: uLSIF, Logistic reg. < KDE < KMM, KLIEP

Computation time of uLSIF and Logistic reg. including cross-validation:





# Covariate Shift: Bias Correction using Density Ratio

training data:  $\{(x_1^{(\text{tr})}, y_1^{(\text{tr})}), \dots, (x_n^{(\text{tr})}, y_n^{(\text{tr})})\}, \{x_1^{(\text{te})}, \dots, x_m^{(\text{te})}\}$

1. Importance estimation:  $x^{(\text{tr})}, x^{(\text{te})} \xrightarrow{\text{estimate}} \hat{w}(x) \cong \frac{p^{(\text{te})}(x)}{p^{(\text{tr})}(x)}$

2. Weighted least-square estimation:

$$\min_{\theta} \frac{1}{n} \sum_{i=1}^n \hat{w}(x_i^{(\text{tr})}) (y_i^{(\text{tr})} - f(x_i^{(\text{tr})}; \theta))^2 + \gamma \|\theta\|_2^2, \quad f(x; \theta) = \sum_{\ell=1}^t \theta_{\ell} K_h(x, m_{\ell}).$$

(Hyper-parameters  $h, \gamma$  are chosen by *importance weighted CV*. cf. Sugiyama et al., nips'07)

Note:

$$\begin{aligned} \frac{1}{n} \sum_{i=1}^n \hat{w}(x_i^{(\text{tr})}) (y_i^{(\text{tr})} - f(x_i^{(\text{tr})}))^2 &\cong \int \frac{p^{(\text{te})}(x)}{p^{(\text{tr})}(x)} (y - f(x))^2 \cdot p(y|x) p^{(\text{tr})}(x) dx \\ &= \int (y - f(x))^2 \cdot p(y|x) p^{(\text{te})}(x) dx \end{aligned}$$

Minimization of weighted square error will provide an asymptotically unbiased estimator of  $f^*(x) = E[Y|X]$  under  $p^{(\text{te})}(x)$ .

# Covariate Shift: Numerical Results

Data set  $\{(x_i, y_i)\}_{i=1}^n$ :  $x_i = (x_i^{(1)}, \dots, x_i^{(d)}) \in [0, 1]^d$  (normalized).

$\{(x_i, y_i)\}_{i=1}^n \xrightarrow{\text{random}} (x_k, y_k)$ ; accepted as  $(x^{(\text{te})}, y^{(\text{te})})$  with probability  $\min\{1, (x_k^{(c)})^2\}$ .

The coordinate  $c$  is randomly determined and fixed in each trial.

$(x^{(\text{tr})}, y^{(\text{tr})})$ : uniformly sampled from the rest.

Data	Uniform	KDE (CV)	KMM (med)	LogReg (CV)	KLIEP (CV)	uLSIF (CV)
kin-8fh	1.00(0.34)	1.22(0.52)	1.55(0.39)	1.31(0.39)	<b>0.95(0.31)</b>	<b>1.02(0.33)</b>
kin-8fm	1.00(0.39)	1.12(0.57)	1.84(0.58)	1.38(0.57)	<b>0.86(0.35)</b>	<b>0.88(0.39)</b>
kin-8nh	<b>1.00(0.26)</b>	1.09(0.20)	1.19(0.29)	1.09(0.19)	<b>0.99(0.22)</b>	<b>1.02(0.18)</b>
kin-8nm	<b>1.00(0.30)</b>	1.14(0.26)	1.20(0.20)	1.12(0.21)	<b>0.97(0.25)</b>	1.04(0.25)
abalone	<b>1.00(0.50)</b>	1.02(0.41)	<b>0.91(0.38)</b>	<b>0.97(0.49)</b>	<b>0.94(0.67)</b>	<b>0.96(0.61)</b>
image	<b>1.00(0.51)</b>	0.98(0.45)	1.08(0.54)	<b>0.98(0.46)</b>	<b>0.94(0.44)</b>	<b>0.98(0.47)</b>
ringnorm	1.00(0.04)	0.87(0.04)	<b>0.87(0.04)</b>	0.95(0.08)	0.99(0.06)	0.91(0.08)
twonorm	1.00(0.58)	1.16(0.71)	<b>0.94(0.57)</b>	<b>0.91(0.61)</b>	<b>0.91(0.52)</b>	<b>0.88(0.57)</b>
waveform	1.00(0.45)	1.05(0.47)	0.98(0.31)	<b>0.93(0.32)</b>	<b>0.93(0.34)</b>	<b>0.92(0.32)</b>
<b>Average</b>	<b>1.00</b>	<b>1.07</b>	<b>1.17</b>	<b>1.07</b>	<b>0.94</b>	<b>0.96</b>
Comp. time	—	<b>0.82</b>	<b>3.50</b>	<b>3.27</b>	<b>2.23</b>	<b>1.00</b>

(average on 100 trials. Wilcoxon signed rank test at the significance level 1%)

# Application 2: Outlier Detection

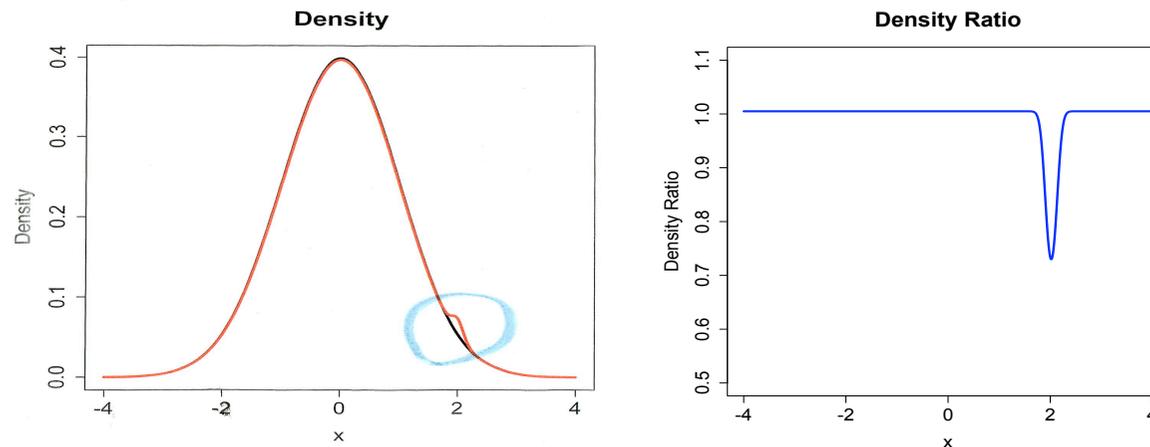
Identify irregular samples in an evaluation dataset.

$$\begin{aligned} \text{Model dataset (no outliers)} : x_1^{(te)}, \dots, x_m^{(te)} &\sim p^{(te)}(x) \\ \text{Evaluation dataset} : x_1^{(tr)}, \dots, x_n^{(tr)} &\sim p^{(tr)}(x) \end{aligned} \xrightarrow{\text{estimate}} w(x) = \frac{p^{(te)}(x)}{p^{(tr)}(x)}$$

- For almost all samples in evaluation data:  $w(x^{(tr)}) = \frac{p^{(te)}(x^{(tr)})}{p^{(tr)}(x^{(tr)})} \cong 1$ .
- On outlying samples in evaluation data:

$$w(x^{(tr)}) = \frac{p^{(te)}(x^{(tr)})}{p^{(tr)}(x^{(tr)})} < 1, \quad (p^{(te)}(x^{(tr)}) < p^{(tr)}(x^{(tr)}))$$

- $w(x)$  can be used as a score of *outlyingness* comparing to model dataset.

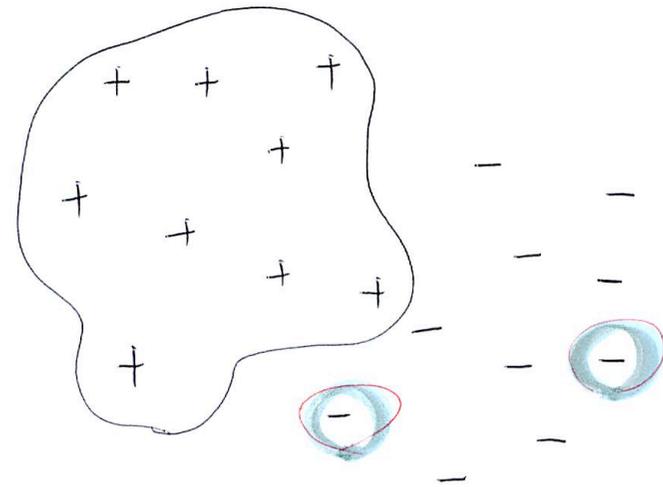


Applications: Intrusion detection in network systems, Topic detection in news documents.

# Outlier Detection: Numerical Experiments

- Benchmark datasets for binary classification problems
  - Model data: training samples with positive label.
  - Evaluation data: test samples with positive label
    - +  $\rho\%$  negative labeled test samples ( $\rho = 1, 2, 5$ )

- Negative labeled samples are randomly chosen from test data set.
- On each dataset, results are averaged on 20 trials.



# Outlier Detection: Results on Benchmark Data

Evaluation: Area under the curve (AUC) of ROC curve: larger is better.

Data		uLSIF (CV)	KLIEP (CV)	LogReg (CV)	KMM (med)	OSVM (med)	LOF			KDE' (CV)	
Name	$\rho$						$k = 5$	$k = 30$	$k = 50$		
banana	1	0.851	0.815	0.447	0.578	0.360	0.838	0.915	0.919	0.934	
	2	0.858	0.824	0.428	0.644	0.412	0.813	0.918	0.920	0.927	
	5	0.869	0.851	0.435	0.761	0.467	0.786	0.907	0.909	0.923	
b-cancer	1	0.463	0.480	0.627	0.576	0.508	0.546	0.488	0.463	0.400	
	2	0.463	0.480	0.627	0.576	0.506	0.521	0.445	0.428	0.400	
	5	0.463	0.480	0.627	0.576	0.498	0.549	0.480	0.452	0.400	
diabetes	1	0.558	0.615	0.599	0.574	0.563	0.513	0.403	0.390	0.425	
	2	0.558	0.615	0.599	0.574	0.563	0.526	0.453	0.434	0.425	
	5	0.532	0.590	0.636	0.547	0.545	0.536	0.461	0.447	0.435	
f-solar	1	0.416	0.485	0.438	0.494	0.522	0.480	0.441	0.385	0.378	
	2	0.426	0.456	0.432	0.480	0.550	0.442	0.406	0.343	0.374	
	5	0.442	0.479	0.432	0.532	0.576	0.455	0.417	0.370	0.346	
		⋮	other 8 datasets				⋮				⋮
Average		<b>0.661</b>	<b>0.685</b>	0.530	0.608	0.596	0.594	0.629	0.622	0.623	
Comp. time		<b>1.00</b>	11.7	5.35	751	12.4	85.5			8.70	

# Conclusion

- A new estimator for density ratio has been proposed
  - Analytic computation of estimator and LOOCV
- Applications: covariate-shift adaptation and outlier detection

## Future works

- Explore various possible applications of density ratio:
  - feature selection, independent component analysis, dimensionality reduction, . . . . .