

Fisher Discriminant Analysis with Kernels

- by Mika, Rätsch, Weston, Schölkopf, Müller
- presented by Boley.

Discriminate between two classes

- Need to identify good set of features
- PCA: unsupervised algorithm to reduce reconstruction error
- Better to take advantage of label info
- Classical approaches: bayes classifier - requires assumptions on data distribution within each class
- Often: assume Gaussian distribution within each class
 - leads to quadratic or linear discriminants, like Fisher

This work

- Authors propose kernel idea used in SVMs, K-PCA.
- Use in supervised Fisher's Discriminant
- Result often competitive with K SVMs.
- Dot-product in kernel space \rightarrow closed form solution

Classical Fisher Linear Discriminant

- samples from two classes: $X_1 = [\mathbf{x}_1, \dots, \mathbf{x}_{\ell_1}]$, $X_2 = [\mathbf{x}_{\ell_1+1}, \dots, \mathbf{x}_{\ell_1+\ell_2}]$, with $\ell = \ell_1 + \ell_2$.
- Fisher's discriminant projects all the data onto a direction \mathbf{w} maximizing the separation of the means along the projection while minimizing the scatter within each class

$$\max J(\mathbf{w}) = \frac{\mathbf{w}^T S_B \mathbf{w}}{\mathbf{w}^T S_W \mathbf{w}}$$

where

S_B	$=$	$(\mathbf{m}_1 - \mathbf{m}_2)(\mathbf{m}_1 - \mathbf{m}_2)^T$	between cluster scatter
S_W	$=$	$\sum_{i=1,2} \sum_{\mathbf{x} \in X_i} (\mathbf{x} - \mathbf{m}_i)(\mathbf{x} - \mathbf{m}_i)^T$	within class scatter
\mathbf{m}_i	$=$	$\frac{1}{\ell_i} \sum_{\mathbf{x} \in X_i} \mathbf{x}$	class mean
\mathbf{m}	$=$	$\frac{1}{\ell} \sum_{\mathbf{x}} \mathbf{x} = \frac{\ell_1}{\ell} \mathbf{m}_1 + \frac{\ell_2}{\ell} \mathbf{m}_2$	global mean

Statistical Motivation - Bayes

- Optimal Bayes assigns class based on maximum a-posteriori probability
- Simplifying assumption: each class has a normal distribution
- Measures Mahalanobis distance of a sample to class center
- Result is a quadratic separator
- With a single common Covariance matrix \rightarrow linear separator
- linear separator advantage: robust against noise
- Direction of separator aligned with direction of maximal variance within each class
- Linear separator \leftrightarrow Fisher's w .
- Crucial: have enough samples to get good estimate of Covariance.

Fisher's discriminant in feature space

- Linear discriminant is not rich enough
- Want to keep robustness and statistical foundation while allowing richer separators
- Answer: use high-dimensional feature space \mathcal{F}
- Map $\mathbf{x} \mapsto \hat{\mathbf{x}} = \phi(\mathbf{x}) \in \mathcal{F}$.
- Fisher's Disc. is now:

$$\max J(\mathbf{w}) = \frac{\mathbf{w}^T \hat{S}_B \mathbf{w}}{\mathbf{w}^T \hat{S}_W \mathbf{w}}$$

where

\hat{S}_B	$=$	$(\hat{\mathbf{m}}_1 - \hat{\mathbf{m}}_2)(\hat{\mathbf{m}}_1 - \hat{\mathbf{m}}_2)^T$	between cluster scatter
\hat{S}_W	$=$	$\sum_{i=1,2} \sum_{\hat{\mathbf{x}} \in \hat{X}_i} (\hat{\mathbf{x}} - \hat{\mathbf{m}}_i)(\hat{\mathbf{x}} - \hat{\mathbf{m}}_i)^T$	within class scatter
$\hat{\mathbf{m}}_i$	$=$	$\frac{1}{\ell_i} \sum_{\hat{\mathbf{x}} \in \hat{X}_i} \hat{\mathbf{x}}$	class mean
$\hat{\mathbf{m}}$	$=$	$\frac{1}{\ell} \sum_{\hat{\mathbf{x}}} \mathbf{x} = \frac{\ell_1}{\ell} \hat{\mathbf{m}}_1 + \frac{\ell_2}{\ell} \hat{\mathbf{m}}_2$	global mean

Kernel Function

- Need to formulate problem in terms of dot-products of input patterns
- Any solution \mathbf{w} must lie in span of training samples $\hat{\mathbf{x}}_1, \dots, \hat{\mathbf{x}}_\ell$ in \mathcal{F} .
- $\mathbf{w} = \sum_1^\ell \alpha_j \hat{\mathbf{x}}_j = \sum_1^\ell \alpha_j \phi(\mathbf{x}_j)$.

- Inner Product with mean: $\mathbf{w}^T \hat{\mathbf{m}}_i = \sum_{j=1}^{\ell} \alpha_j \underbrace{\frac{1}{\ell_i} \sum_{\mathbf{x} \in X_i} k(\mathbf{x}_j, \mathbf{x})}_{(\mathbf{M}_i)_j}$.

- Wish to optimize $\max J(\mathbf{w}) = \mathbf{w}^T \hat{S}_B \mathbf{w} / \mathbf{w}^T \hat{S}_W \mathbf{w}$

- Numerator: $\mathbf{w}^T \hat{S}_B \mathbf{w} = \alpha^T \underbrace{(\mathbf{M}_1 - \mathbf{M}_2)(\mathbf{M}_1 - \mathbf{M}_2)^T}_M \alpha$

- Here \mathbf{M}_i is the ℓ -vector of weighted row sums of the kernel matrix $K = \{K_{ij}\} = \{k(\mathbf{x}_i, \mathbf{x}_j)\}_{i,j=1,\dots,\ell}$.

Kernel Function 2

- Wish to optimize $\max J(\mathbf{w}) = \mathbf{w}^T \hat{S}_B \mathbf{w} / \mathbf{w}^T \hat{S}_W \mathbf{w}$
- Denominator: $\mathbf{w}^T \hat{S}_W \mathbf{w} = \underbrace{\alpha^T (K_1(I - \mathbf{1}_{\ell_1})K_1^T) + (K_2(I - \mathbf{1}_{\ell_2})K_2^T)}_N \alpha$

where $K_1 = \{(K_1)_{ij}\} = \{k(\mathbf{x}_i, \mathbf{x}_j)\}_{i=1, \dots, \ell}^{j=1, \dots, \ell_1}$ ($\ell \times \ell_1$ matrix)
 $K_2 = \{(K_2)_{ij}\} = \{k(\mathbf{x}_i, \mathbf{x}_j)\}_{i=1, \dots, \ell}^{j=1, \dots, \ell_2}$ ($\ell \times \ell_2$ matrix)
 $K = (K_1, K_2)$.

Kernel Fisher Discriminant

- KFD is now solved by optimizing

$$\max J(\mathbf{w}) = \frac{\mathbf{w}^T N \mathbf{w}}{\mathbf{w}^T M \mathbf{w}}.$$

- Solve by finding leading eigenvector of $N^{-1}M$ [or better, solve generalized eigenproblem $M\mathbf{w} = \lambda N\mathbf{w}$].
- Project new pattern $\hat{\mathbf{x}} = \phi(\mathbf{x})$ onto \mathbf{w} by

$$\langle \mathbf{w}, \phi(\mathbf{x}) \rangle = \sum_{i=1}^{\ell} \alpha_i k(\mathbf{x}_i, \mathbf{x})$$

Numerical Issues

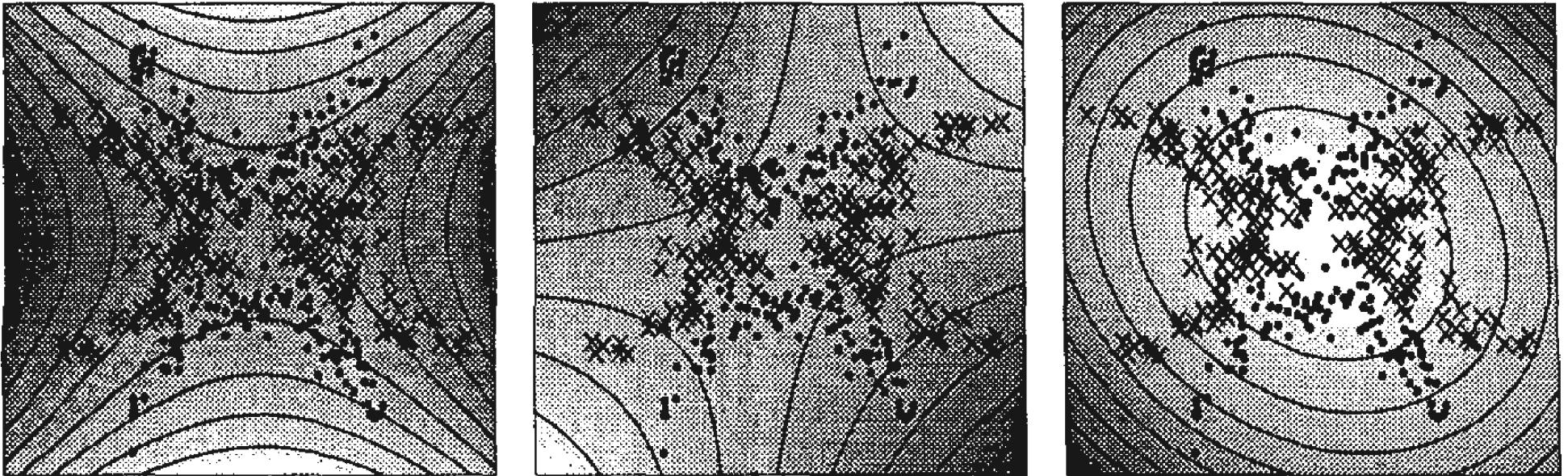
- Estimating ℓ covariance structures from ℓ samples \rightarrow ill-posed.
- N could be singular or badly conditioned
- Need capacity control in \mathcal{F}

Solution

- Replace N with $N_\mu = N + \mu I$.
- Effect: Makes N better conditioned
- Decreases bias in sample-based eigenvalue estimates
- Imposes regularization on $\|\alpha\|^2$, favoring solutions with small expansion coefficients.
- Regularization effect not fully understood.
- Other forms of regularization possible.

Illustration

Figure 1: Comparison of feature found by KFD (left) and those found by Kernel PCA: first (middle) and second (right); details see text.



- KFD: polynomial kernel degree two, regularized with $\mu = 10^{-3}$.
- Two classes (\times 's & \bullet 's), parabolic mirrored around axes.
- Contour lines = level sets
- KFD level sets discriminate classes well
- KPCA less so.

Experiments

- Compare to other state-of-the-art classifiers
- KFD: Kernel Fisher Discriminant with Gaussian kernel
 - Once w obtained, used 1-d linear SVM to classify
- Adaboost
- Regularized Adaboost
- SVM: Support Vector Machine with Gaussian kernel

Data Sets

- Sources: ICI DELVE STATLOG Benchmark data sets
- Treated all as two-class problems
- 100 partitions into training/test sets (about 60%:40%)
- Hyperparameters estimated using 5-fold cross-validation over first 5 realizations
- Table shows average test error & standard deviation over 100 runs

Results

Preliminary Experiment with USPS Digit Data

- Used 3000 training samples
- Compared KFD with KSVM, both with Gaussian kernels
- 10 class error: KFD: 3.7%, KSVM: 4.2%

In General

- Noticed: both KFD & SVM yield optimal hyperplane in \mathcal{F} : often former is better.
- Complexity of SVM classifier is $O(\text{supportvectors})$.
- Complexity of KFD classifier is $O(\text{alltrainingvectors})$.
- Dependence on all training vectors \rightarrow maybe more robust.
- KFD: closed form solution.
Other methods involve a search or an optimization problem.
- Table on next page: **1st place in bold**, *2nd place in italic* (lower is better)

Experiments

Table 1: Comparison between KFD, a single RBF classifier, AdaBoost (AB), regularized AdaBoost (AB_R) and Support Vector Machine (SVM) (see text). Best method in bold face, second best emphasized.

	RBF	AB	AB_R	SVM	KFD
Banana	10.8±0.6	12.3±0.7	<i>10.9±0.4</i>	11.5±0.7	10.8±0.5
B.Cancer	27.6±4.7	30.4±4.7	26.5±4.5	<i>26.0±4.7</i>	25.8±4.6
Diabetes	24.3±1.9	26.5±2.3	23.8±1.8	<i>23.5±1.7</i>	23.2±1.6
German	24.7±2.4	27.5±2.5	24.3±2.1	23.6±2.1	<i>23.7±2.2</i>
Heart	17.6±3.3	20.3±3.4	16.5±3.5	16.0±3.3	<i>16.1±3.4</i>
Image	3.3±0.6	2.7±0.7	2.7±0.6	<i>3.0±0.6</i>	4.8±0.6
Ringnorm	1.7±0.2	1.9±0.3	<i>1.6±0.1</i>	1.7±0.1	1.5±0.1
F.Sonar	34.4±2.0	35.7±1.8	34.2±2.2	32.4±1.8	<i>33.2±1.7</i>
Splice	<i>10.0±1.0</i>	10.1±0.5	9.5±0.7	10.9±0.7	10.5±0.6
Thyroid	4.5±2.1	<i>4.4±2.2</i>	4.6±2.2	4.8±2.2	4.2±2.1
Titanic	23.3±1.3	<i>22.6±1.2</i>	<i>22.6±1.2</i>	22.4±1.0	23.2±2.0
Twonorm	2.9±0.3	3.0±0.3	<i>2.7±0.2</i>	3.0±0.2	2.6±0.2
Waveform	10.7±1.1	10.8±0.6	9.8±0.8	<i>9.9±0.4</i>	<i>9.9±0.4</i>

Conclusions and Discussion

- Fisher's discriminant: standard linear statistical technique, but too limited.
- This is one of many approaches to obtain more general class separability.
- Advantage: closed form solution.
- Flexibility: wide choice of kernels.
- Experimental results: competitive with many other methods.
- Complexity scales with all training samples (not just the difficult ones)

Future Work

- Suitable approximation schemes
- Numerical methods to find a few leading eigenvectors
- Multi-class discriminants
- Generalization bounds.

Novelty Detection

Kernel PCA for Novelty Detection by Heiko Hoffman

- Novelty Detection is a one-class classification problem.
- Use training data to see typical acceptable data.
- Called One-Class because training data contains only acceptable data.
- Test data may be similar to training data or not: objective is to distinguish those that are different.
- Abnormal examples are generally rare.
- Alternate algorithm: One-class SVM: find tightest separator from origin in \mathcal{F} .
- Alternate algorithm: SVDD: Find smallest enclosing sphere in kernel space \mathcal{F} . RBF kernel leads to same as one-class SVM.
- Here we try to generate a simplified model.
- Alternate approaches: • Gaussian Mixture models, • auto-associative multilayer perceptron
 - principal curves and surfaces,All these lead to non-linear (often non-convex) optimization problems.
- Here we use PCA in kernel space to reduce dimensionality.

Method

- Training data are mapped into an infinite-dimensional feature space.
- In this space, kernel PCA extracts the principal components of the data distribution. Eigenvectors $\{\mathbf{v}_\ell\}_{\ell=1}^q$ of \bar{K} with $\bar{K}_{ij} = K_{ij} - \frac{1}{n} \sum_r K_{ir} - \frac{1}{n} \sum_r K_{rj} + \frac{1}{n^2} \sum_{r,s} K_{rs}$ where $K_{ij} = k(\mathbf{x}_i, \mathbf{x}_j)$.

- Potential: $p_S(\mathbf{z}) = \|\phi(\mathbf{z}) - \bar{\phi}\|_2^2 = k(\mathbf{z}, \mathbf{z}) - \frac{2}{n} \sum_{i=1}^n k(\mathbf{z}, \mathbf{x}_i) + \frac{1}{n^2} \sum_{i,j}^n k(\mathbf{x}_i, \mathbf{x}_j)$

- Projection: $f_\ell(\mathbf{z}) = \left\langle \left[\phi(\mathbf{z}) - \frac{1}{n} \sum_{r=1}^n \phi(\mathbf{x}_r) \right], [\mathbf{v}_\ell - \bar{\phi}(\mathbf{x})] \right\rangle$

where $\mathbf{v}_\ell = \ell$ -th eigenvector & $\bar{\phi}(\mathbf{x})$ is center in \mathcal{F} (both linear comb's of $\phi(\mathbf{x}_i)$'s).

- The squared distance to the corresponding principal subspace is the measure for novelty:

$$p(\mathbf{z}) = p_s(\mathbf{z}) - \sum_{i=1}^q f_\ell(\mathbf{z})^2$$

Diagram

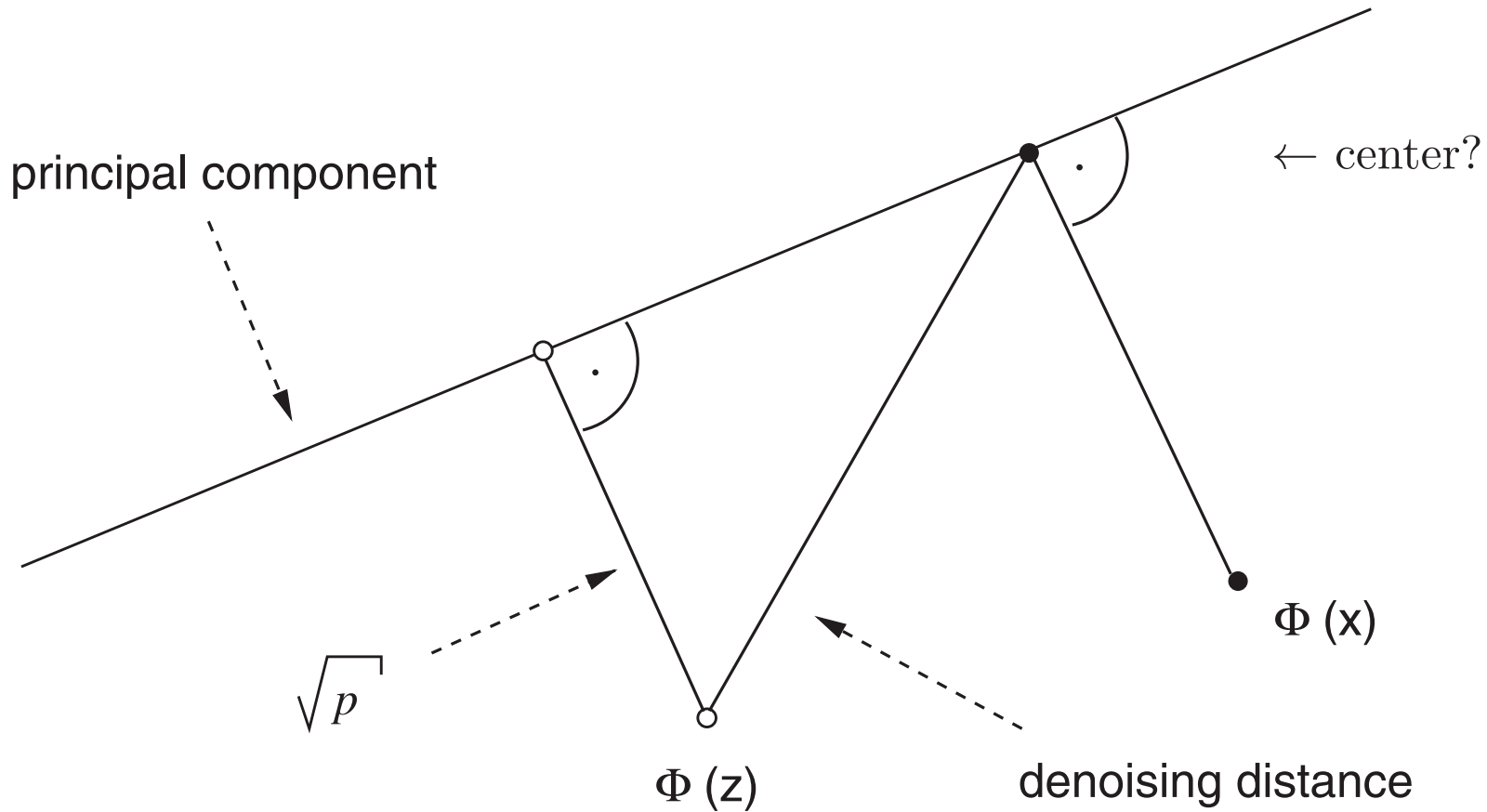
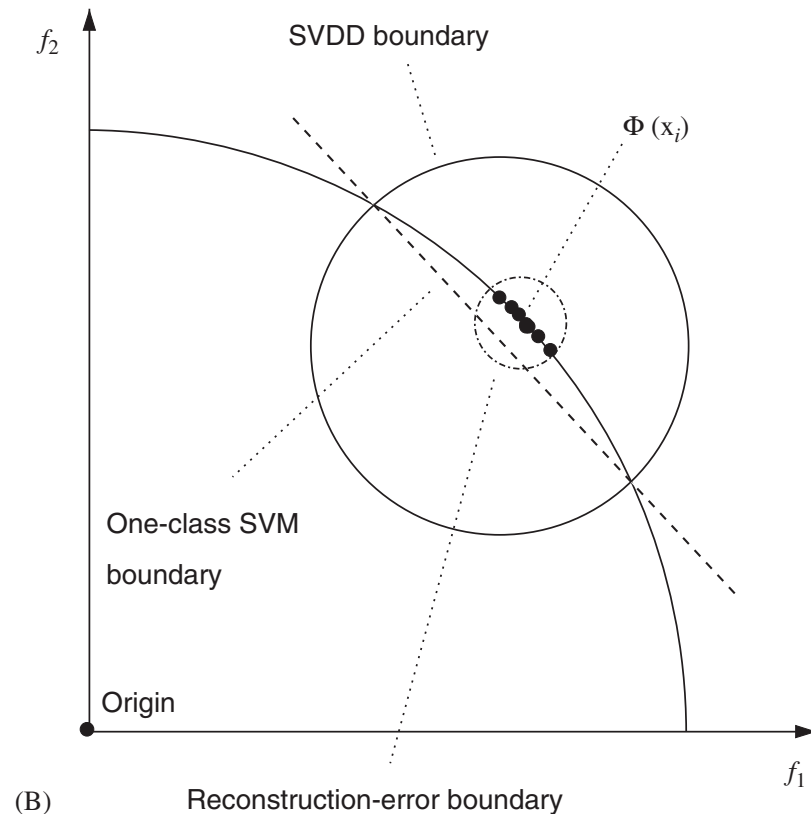
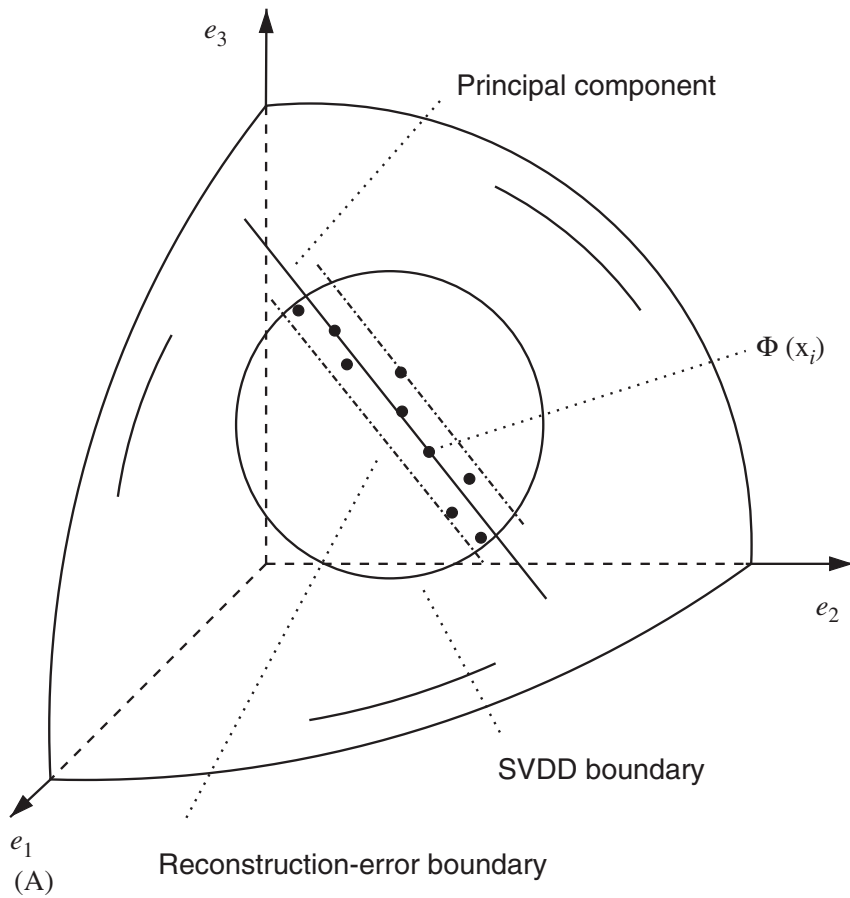


Fig. 12. The difference between the distance to be optimized in denoising and the reconstruction error p .

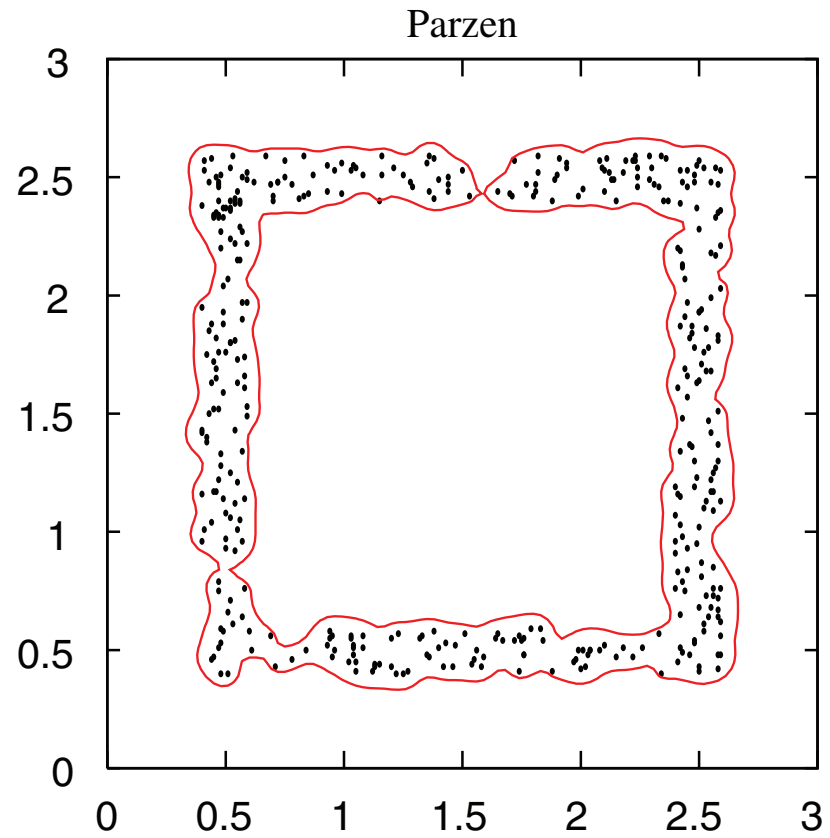
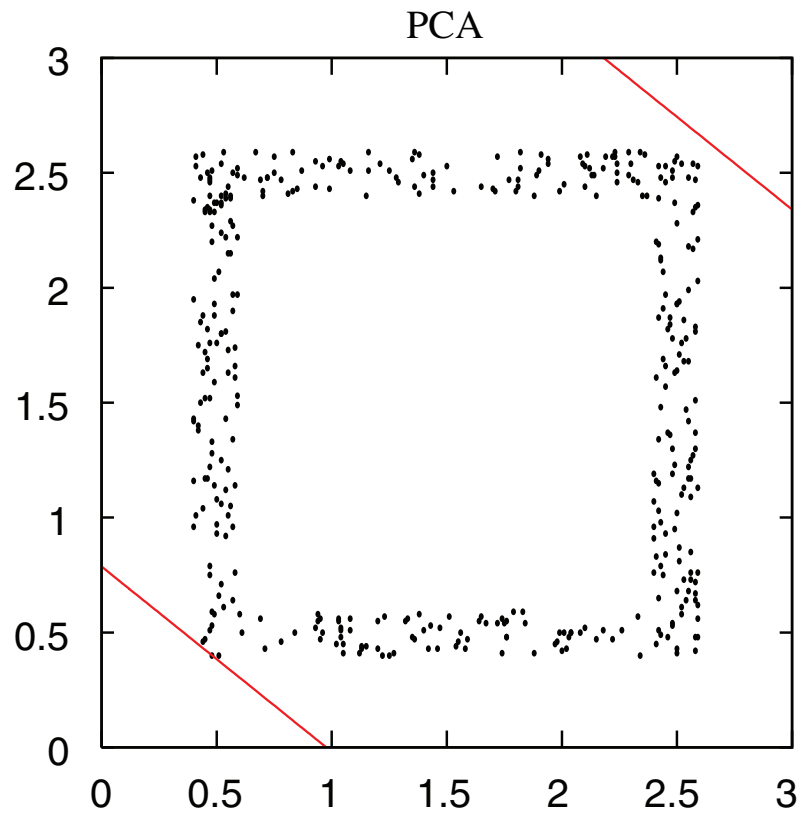
Decision Boundary Sketch

Fig. 1. Decision boundaries in the feature space of an RBF kernel, comparing one-class SVM, SVDD, and the reconstruction error: (A) The boundaries are illustrated in a three-dimensional feature space. All data points $\Phi(\mathbf{x}_i)$ lie on a sphere. (B) Cross-section through the center of the SVDD sphere and orthogonal to the principal component for the situation in A.

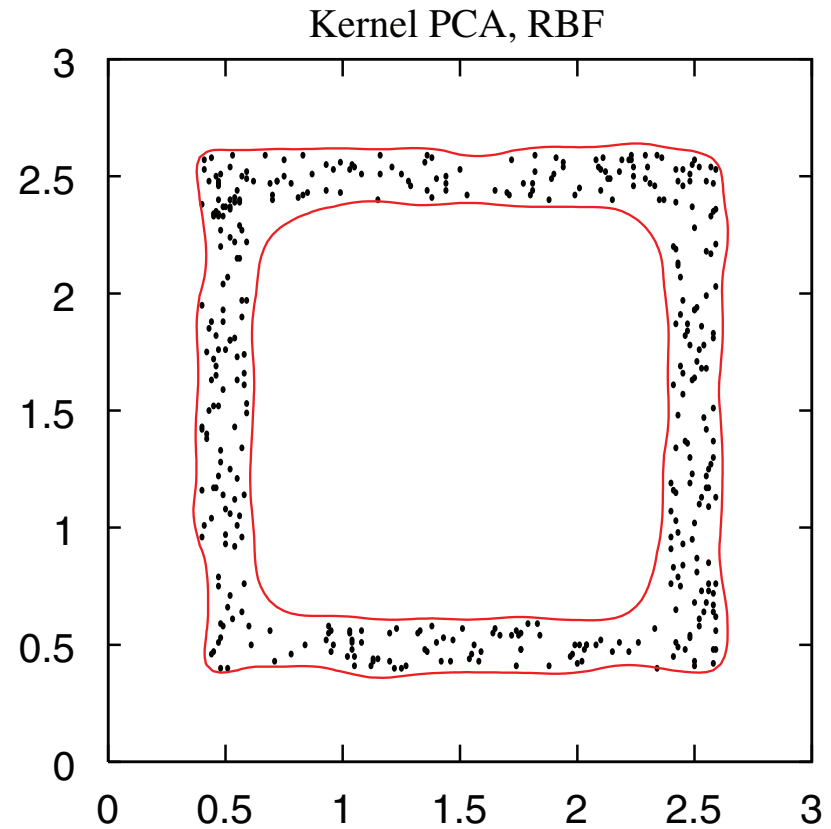
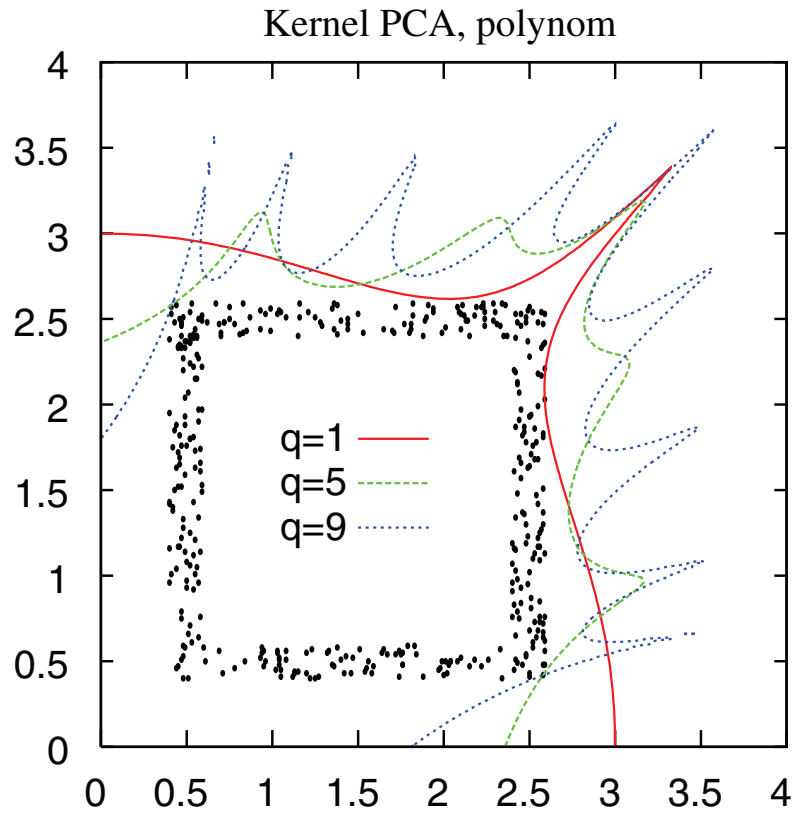
Illustration



Example - classical methods



Example - kernel methods



Ring Square Boundary

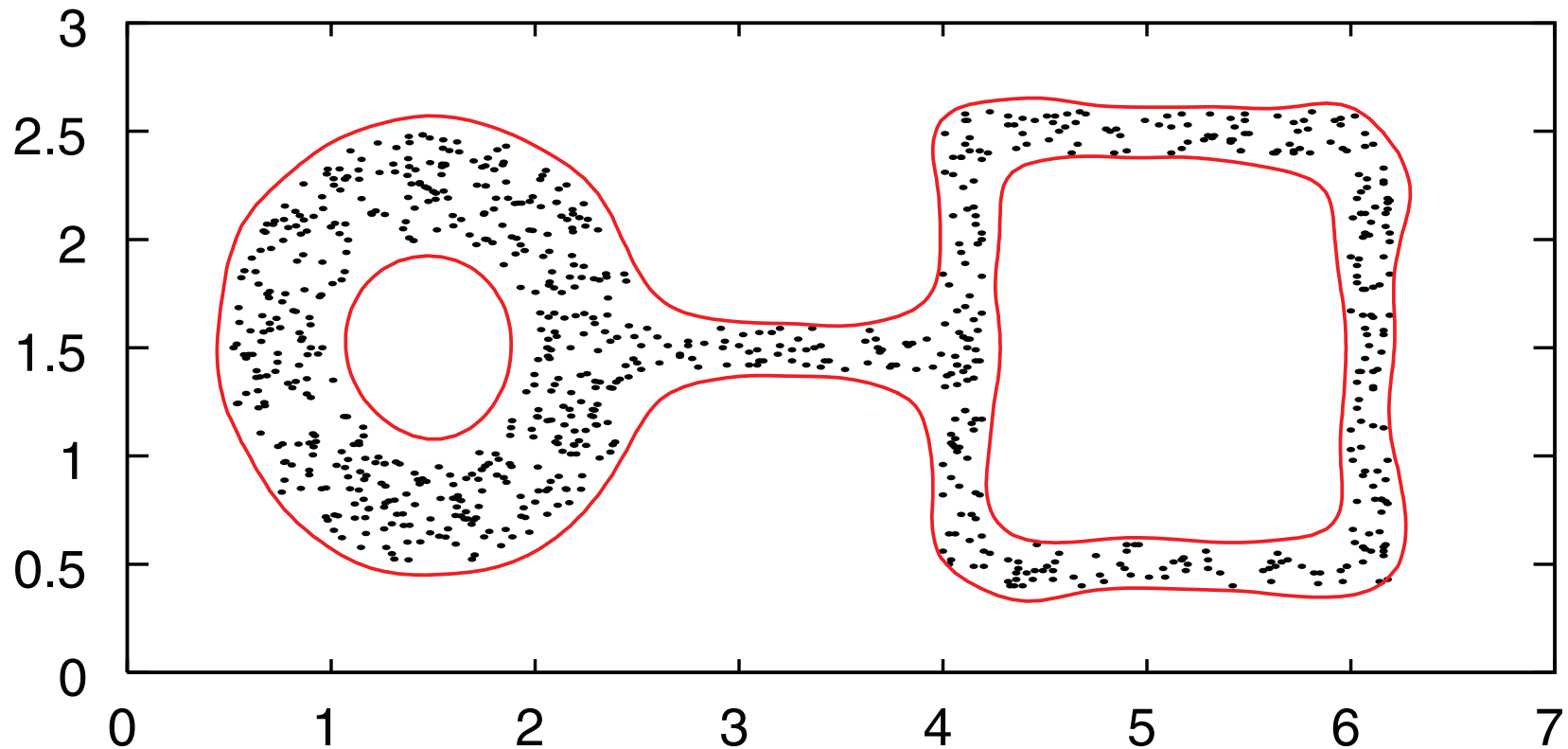


Fig. 3. Decision boundary for the ring-line-square distribution using the reconstruction error in \mathcal{F} with $\sigma = 0.4$ and $q = 40$.

spiral

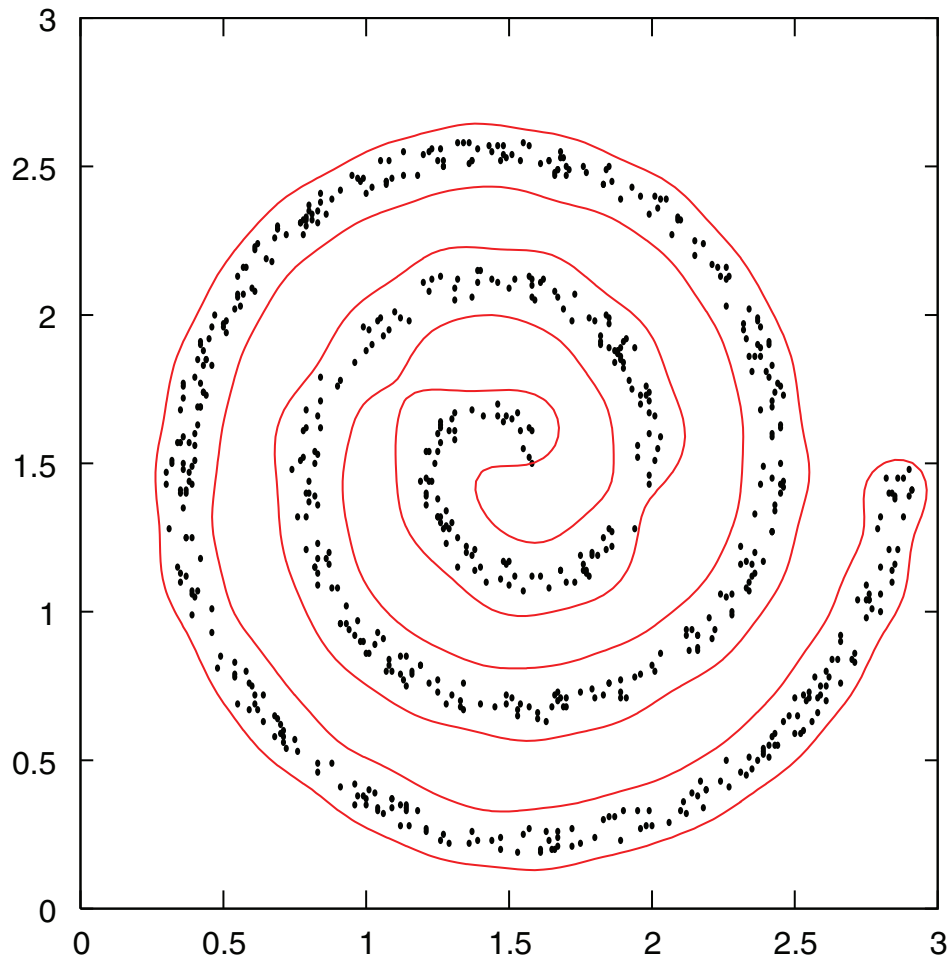
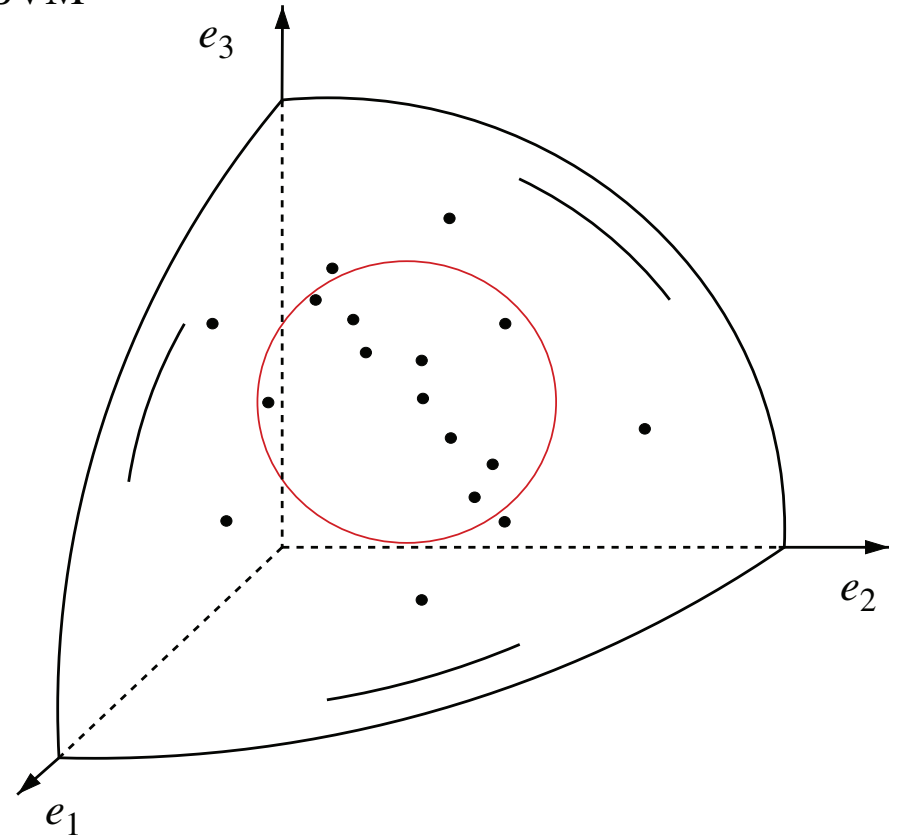
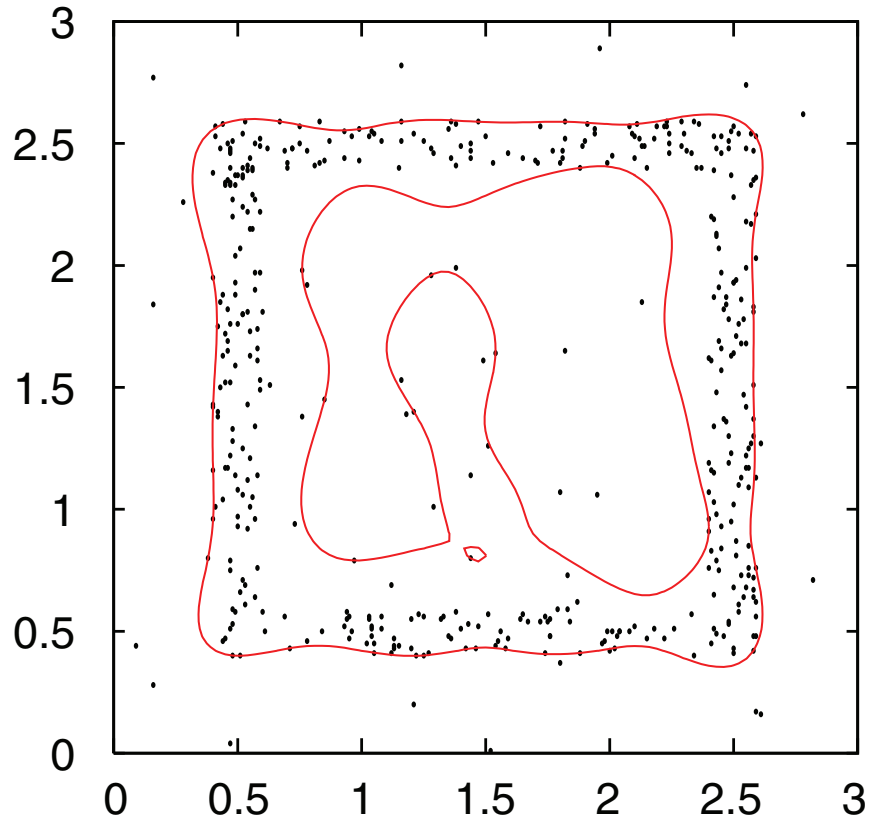


Fig. 4. Decision boundary for the spiral distribution using the reconstruction error in \mathcal{F} with $\sigma = 0.25$ and $q = 40$.

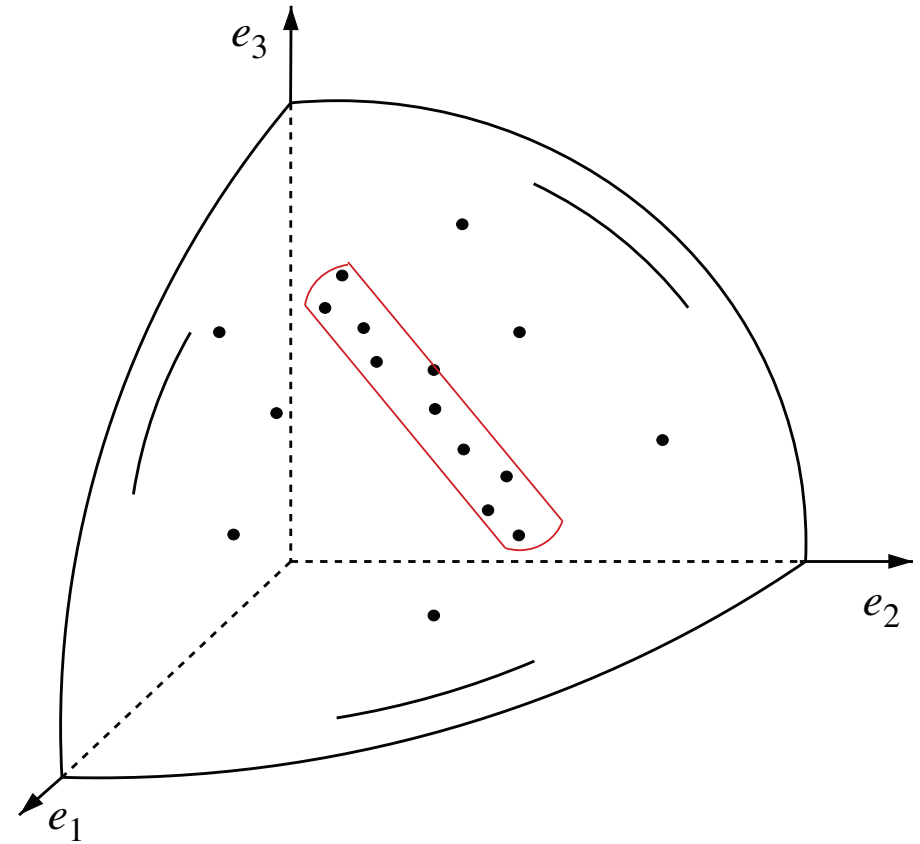
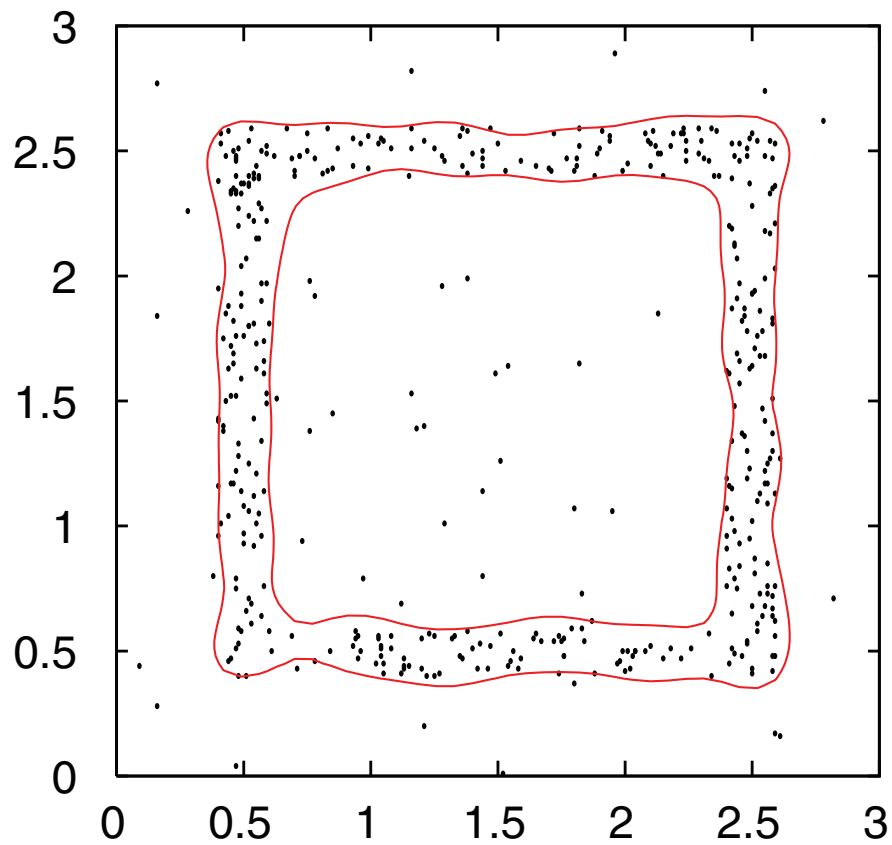
Noisy Data - One-class SVM

One-class SVM



Noisy Data - K-PCA

Kernel PCA



Samples from sine curve plus uniform noise

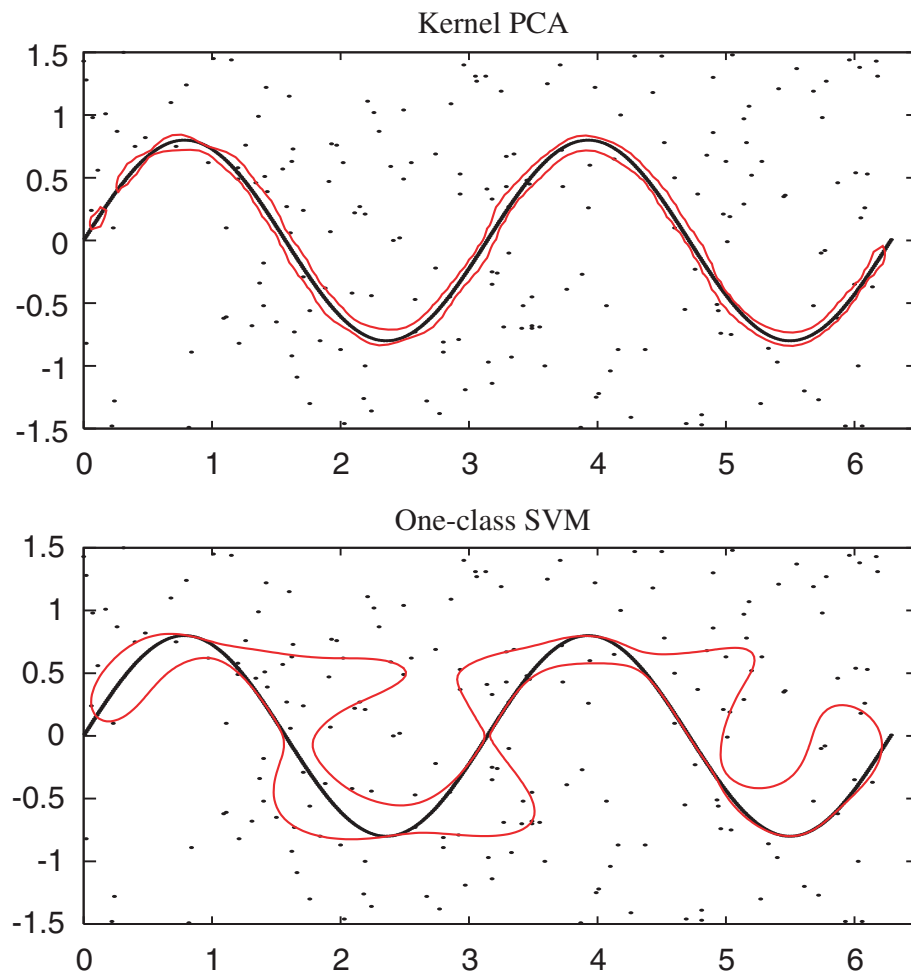
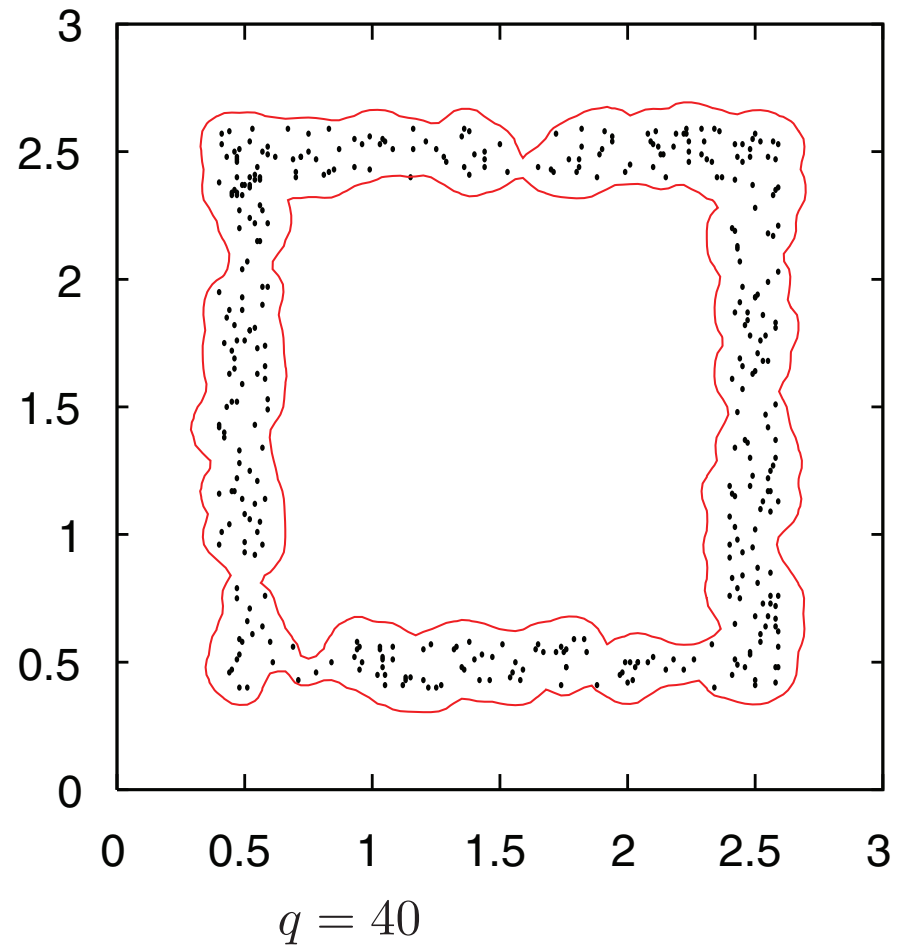
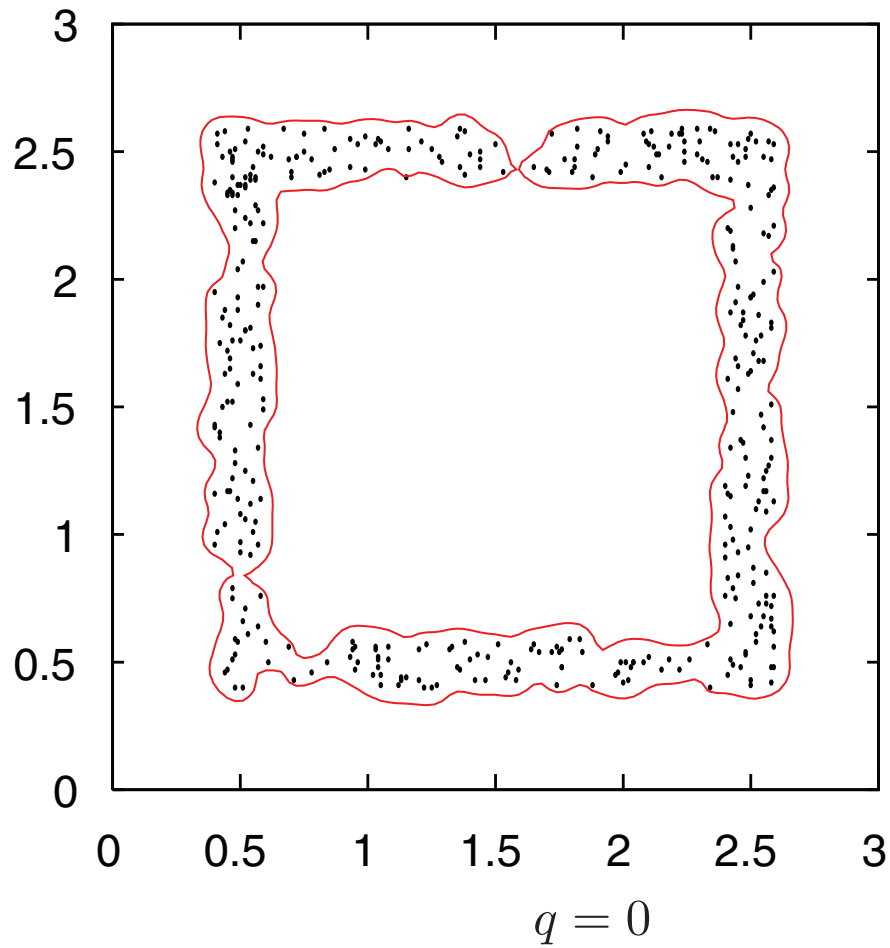
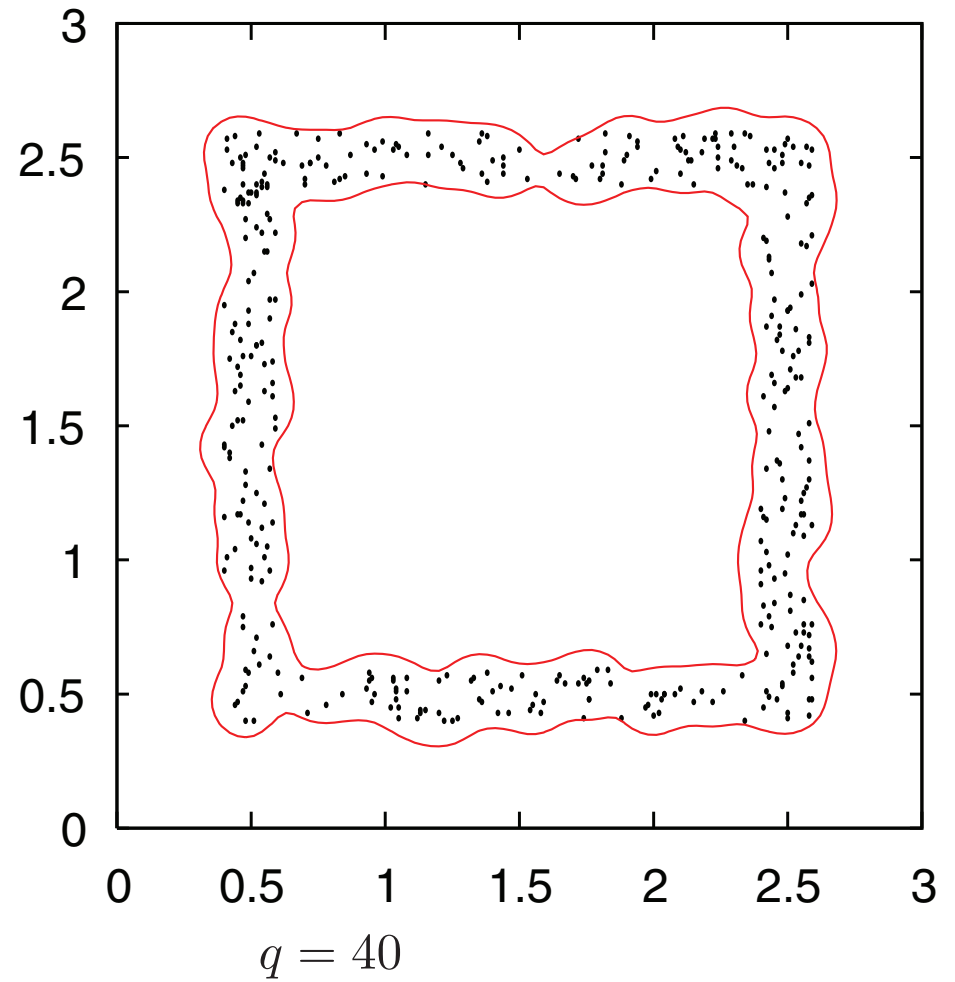
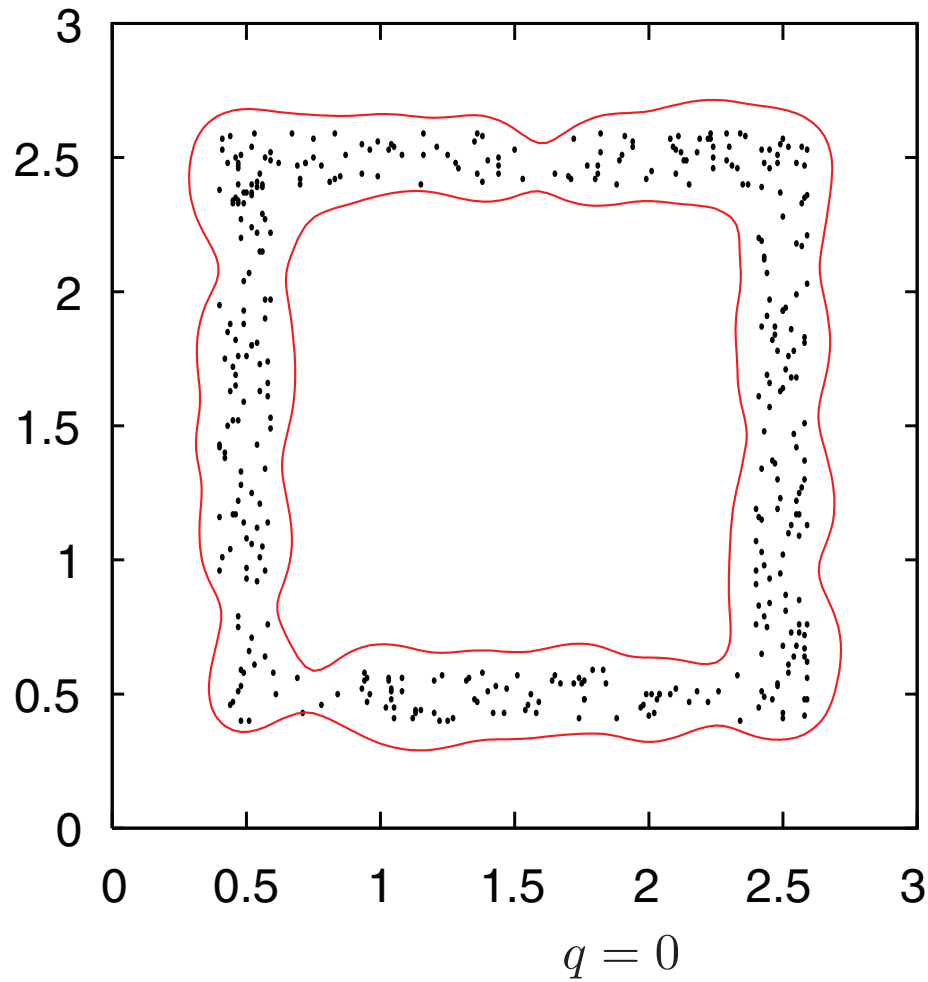


Fig. 6. Decision boundaries for the sine-noise distribution comparing kernel PCA ($\sigma=0.4$, $q=40$) with the one-class SVM ($\sigma=0.489$, $\nu=\frac{2}{7}$).

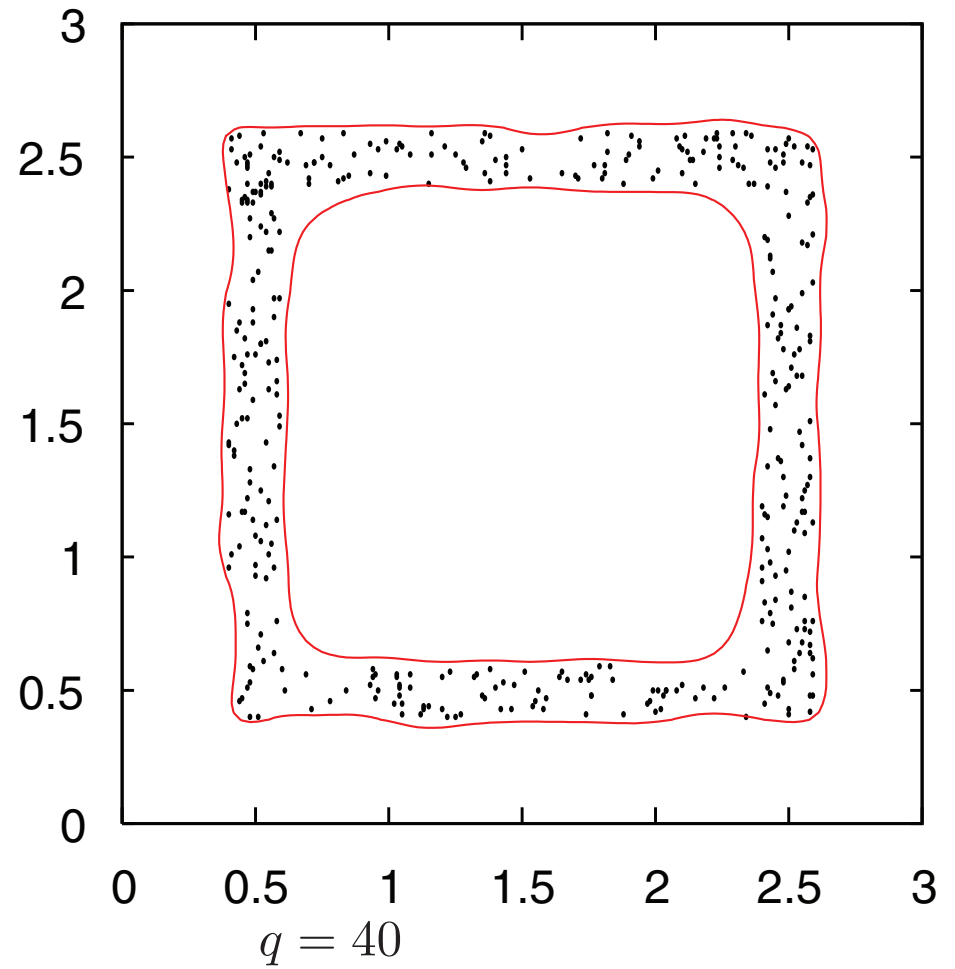
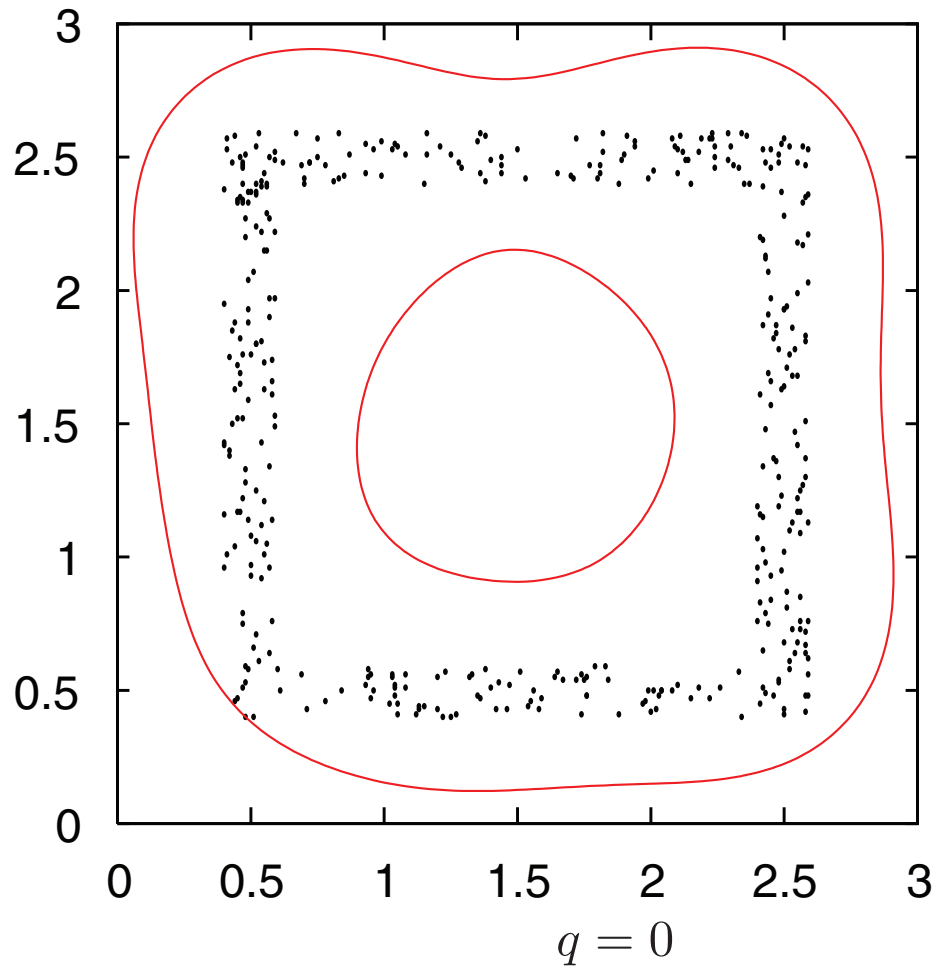
Vary Parameters: $\sigma = .05$



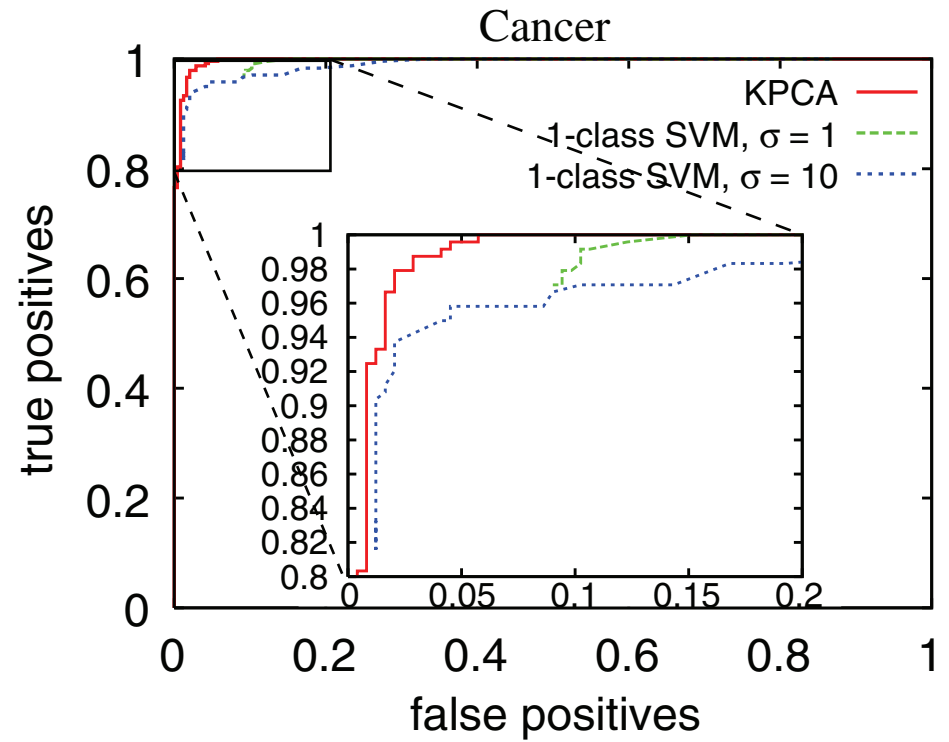
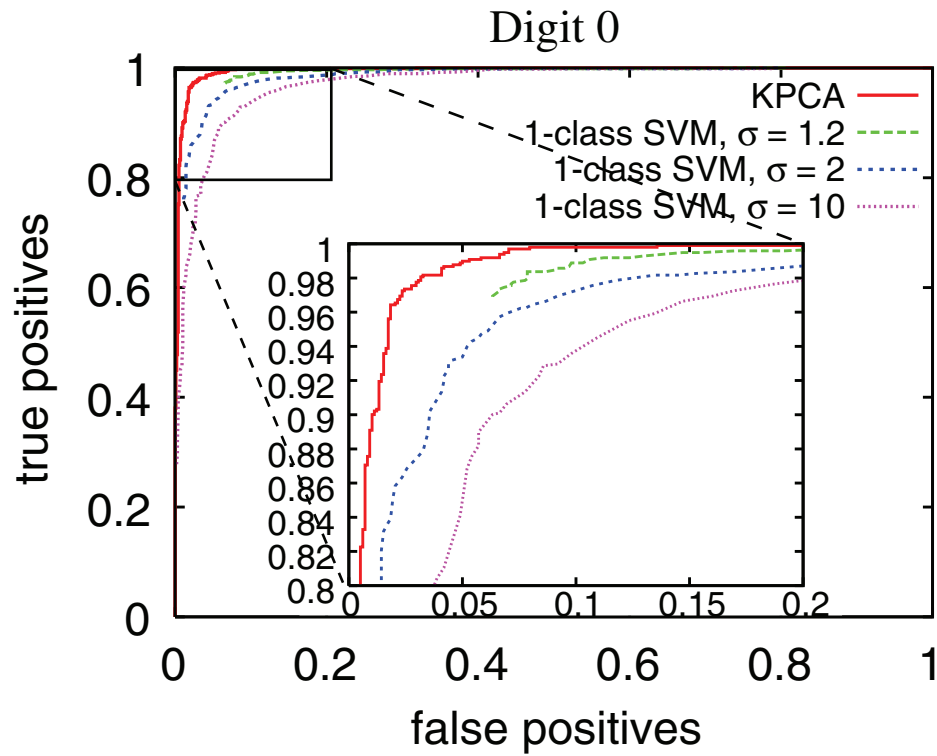
Vary Parameters: $\sigma = .10$



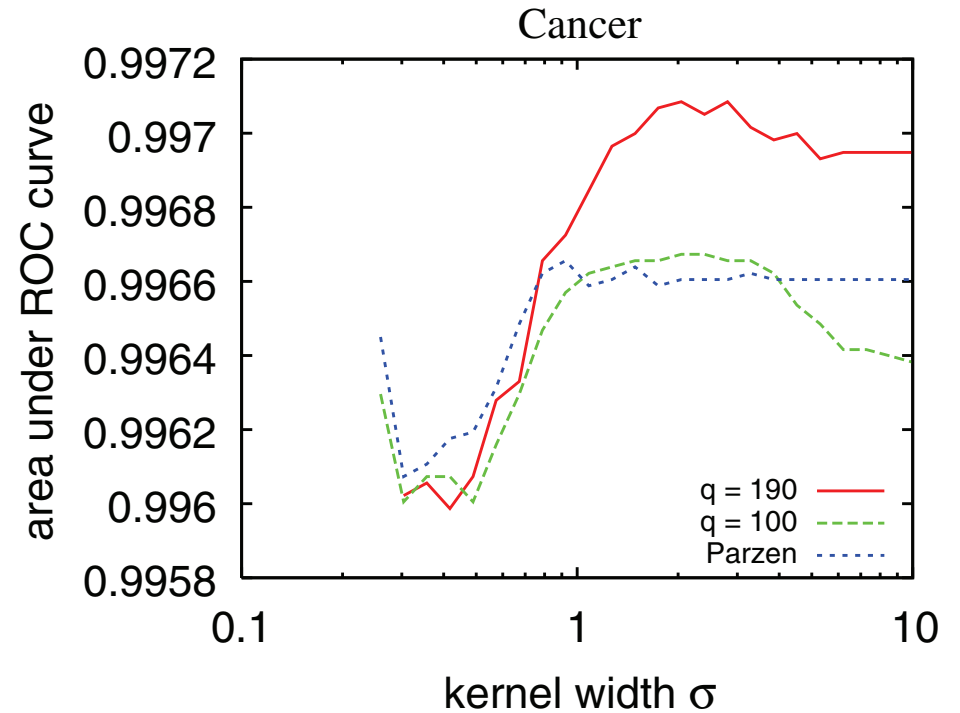
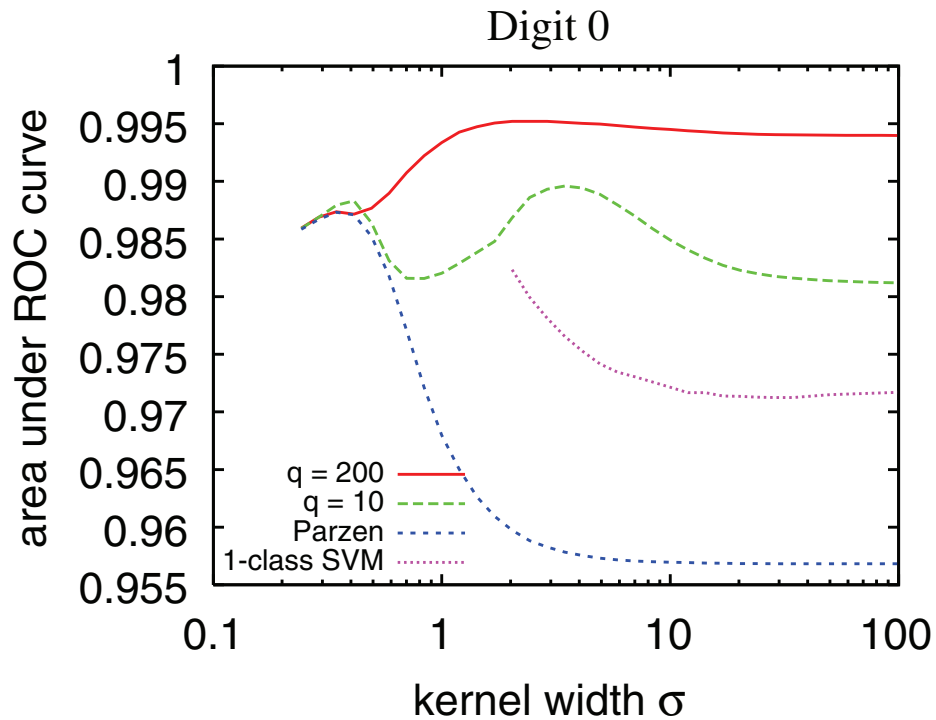
Vary Parameters: $\sigma = .40$



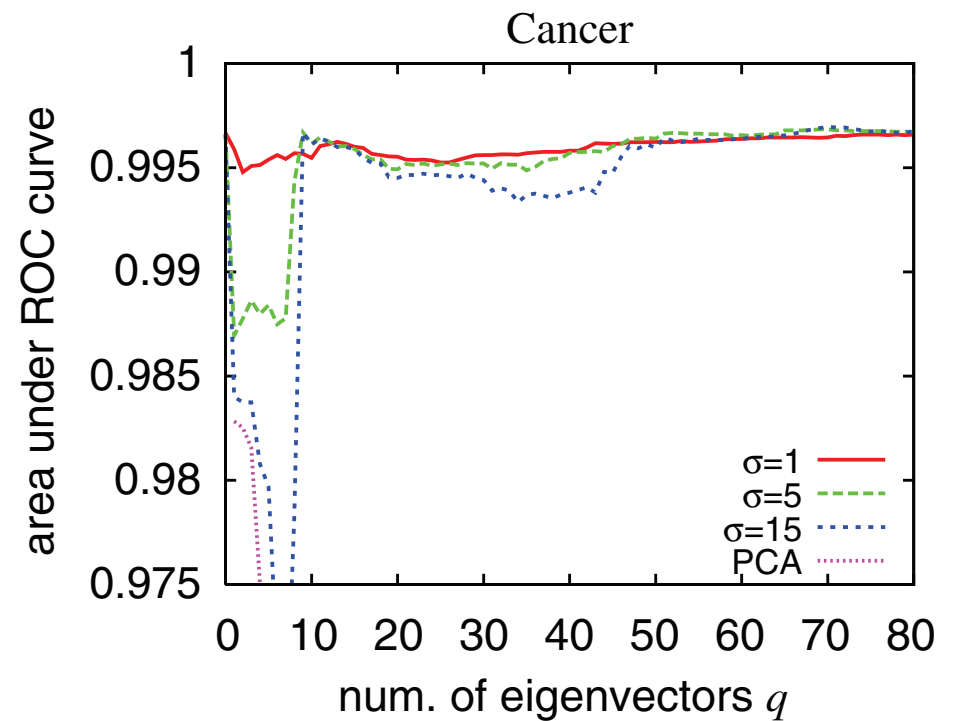
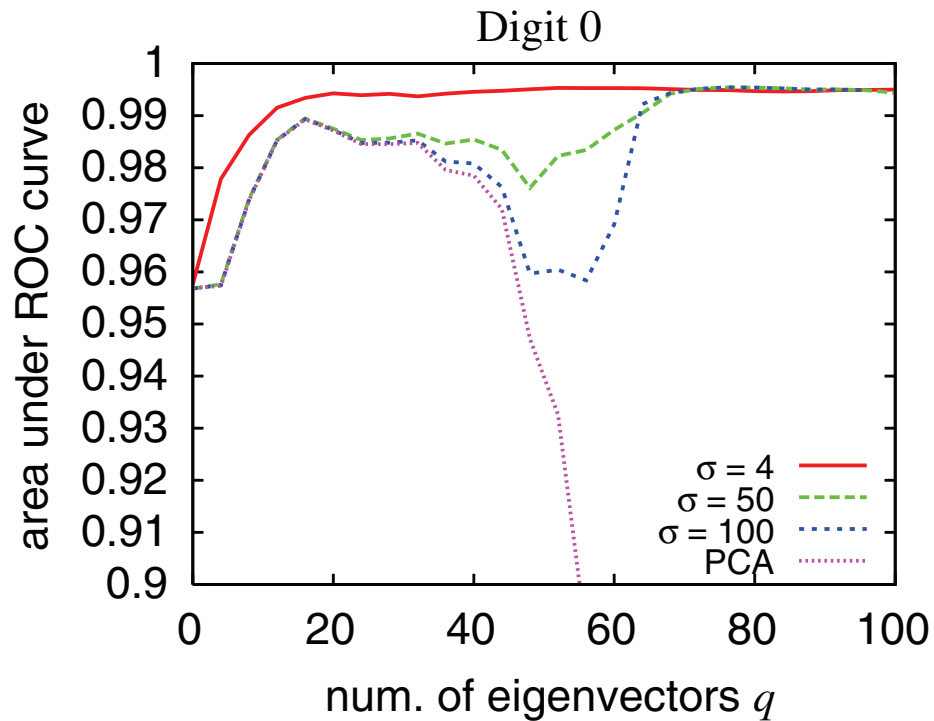
Real Data ROC curves : Classifier



Real Data: vary kernel width



Real Data: vary # eigenvectors



Most Unusual Zero Digits

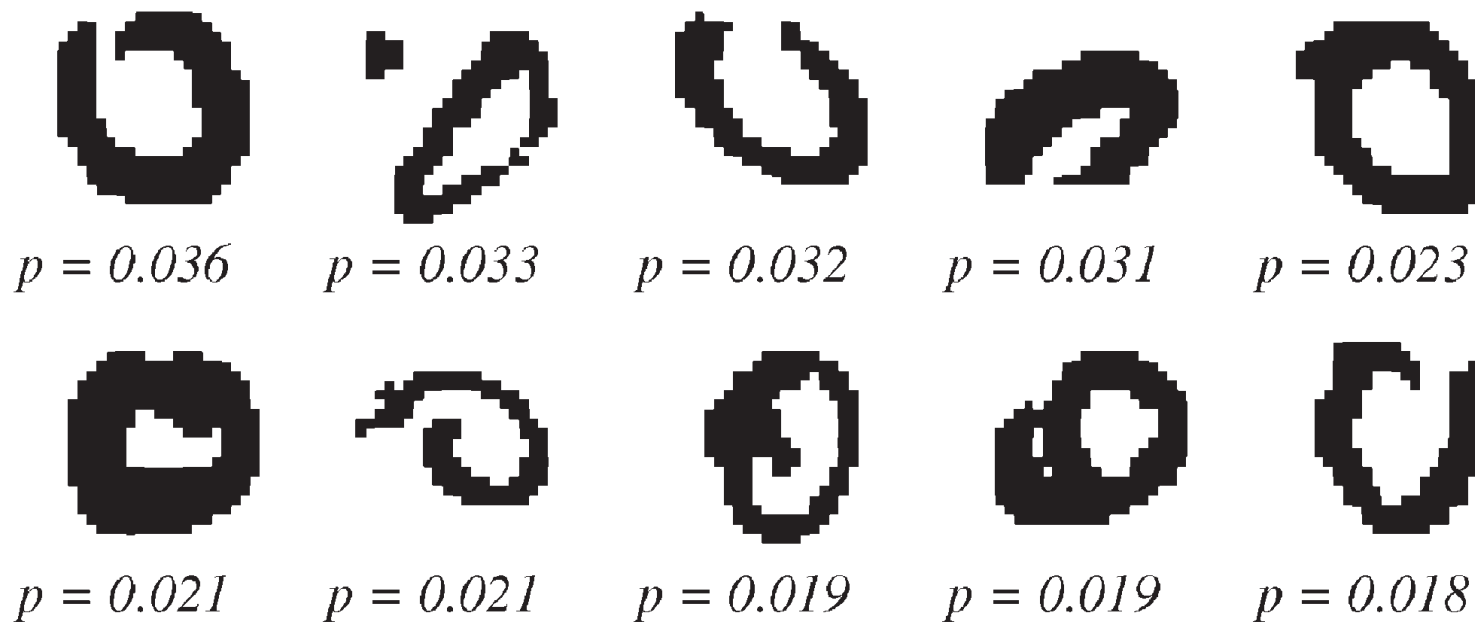


Fig. 11. The 10 most unusual ‘0’ digits from the MNIST test set. The digits are arranged in descending order of their reconstruction error p ($\sigma = 4$, $q = 100$). The figure shows the unprocessed digits of size 28×28 pixels; for novelty detection, however, the processed digits (8×8 pixels) were used.

Diagram

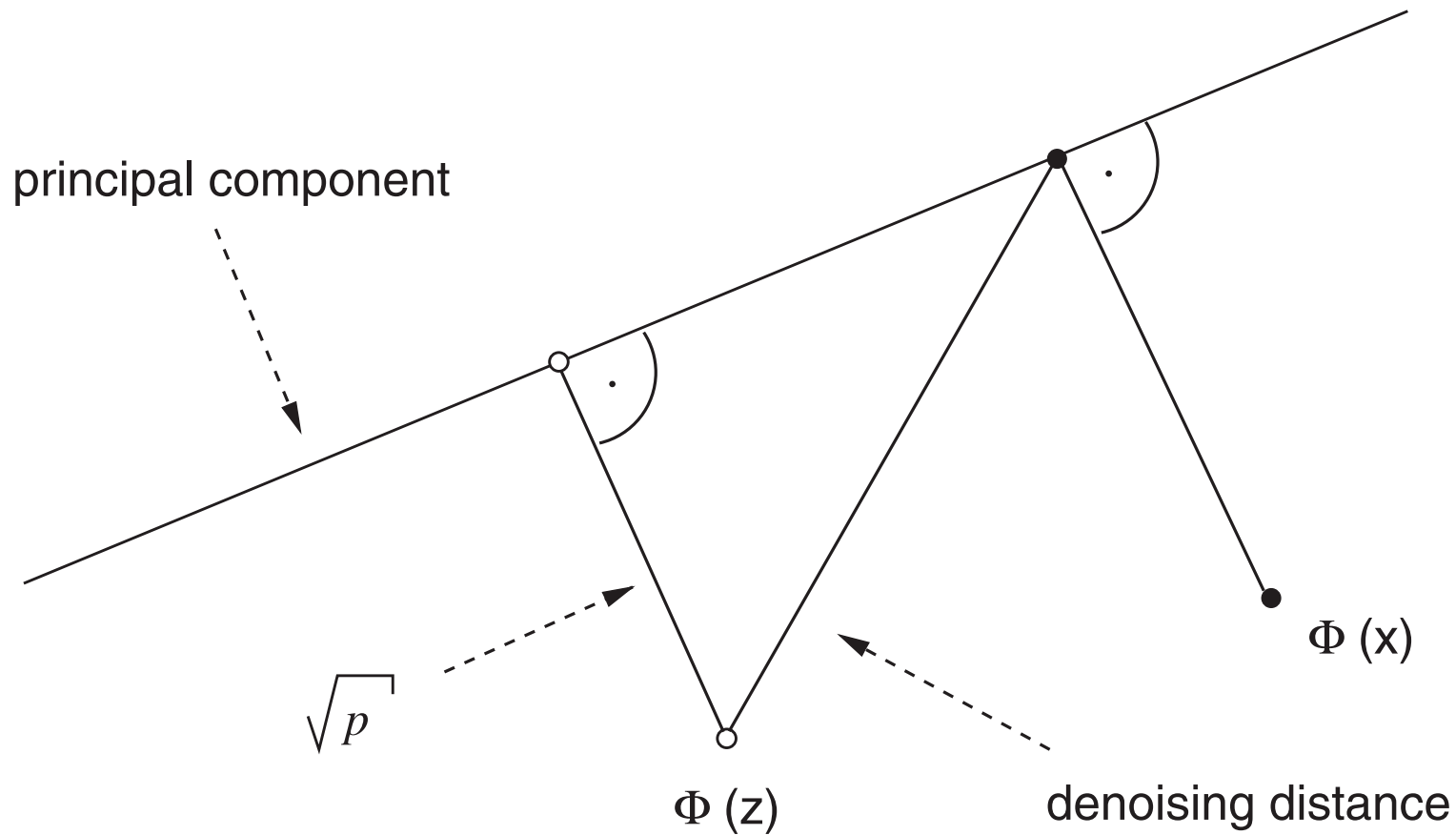


Fig. 12. The difference between the distance to be optimized in denoising and the reconstruction error p .