# 核方法 Kernel Method

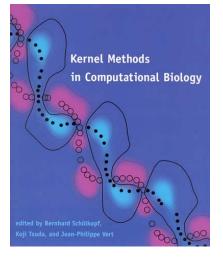
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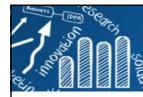






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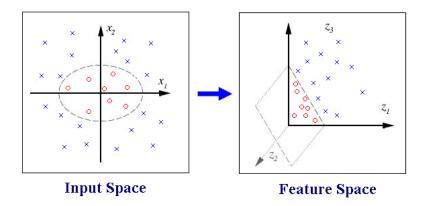




# 核方法 (Kernel Methods)

- Aronszajn (1950) and Parzen (1962) first to employ *kernel methods* in statistics.
- Aizerman et al. (1964) used *positive definite kernels* which was closer to "*kernel trick*", they argue that a *positive definite kernel* is identical to a *dot product* in the feature space.
  - Boser et al (1992), to construct *SVMs*, a generalization of the so-called optimal hyperplane algorithm.

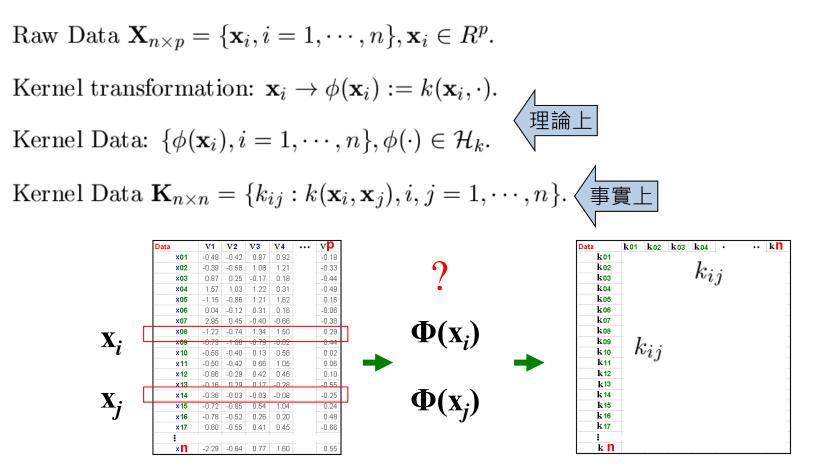
$$\Phi : \mathbb{R}^2 \to \mathbb{R}^3$$
  
(x<sub>1</sub>, x<sub>2</sub>)  $\mapsto$  (z<sub>1</sub>, z<sub>2</sub>, z<sub>3</sub>) := (x<sub>1</sub><sup>2</sup>,  $\sqrt{2} x_1 x_2, x_2^2$ )



- Scholkopf et al (1998) point out that kernels can be used to construct generalization of any algorithm that can be carried out in terms of *dot products*.
- For last 20 years, there have seen a large number of *kernelization* of various algorithms. (PCA, LDA, CCA, PLS,...)



# **Prepare Kernel Data**



- Linear:  $k(x,y) = \langle x,y \rangle$
- Polynomial:  $k(x, y) = (\text{scale} \cdot \langle x, y \rangle + \text{offset})^{\text{degree}}$
- Gaussian Radial Basis Function:  $k(x, y) = \exp\{-\operatorname{scale} \cdot ||x y||^2\}$



Data are not represented individually anymore, but only through a set of pairwise comparisons.

A real-valued comparison function  $k : \mathcal{X} \times \mathcal{X} \to R$  is used, and data set  $\mathbf{X}_{[n \times p]}$  is represented by the  $n \times n$  matrix of pairwise comparisons  $K_{ij} = k(\mathbf{x}_i, \mathbf{x}_j)$ .

- The representation as a square matrix does not depend on the nature of the objects to be analyzed.
- The size of the matrix used to represent a dataset of n objects is always n by n.

**Definition:** a function  $k : \mathcal{X} \times \mathcal{X} \to R$  is called a **positive definite kernel** *iff* it is symmetric, that is,  $k(\mathbf{x}_i, \mathbf{x}_j) = k(\mathbf{x}_j, \mathbf{x}_i)$  for any two objects  $\mathbf{x}_i, \mathbf{x}_j$  in  $\mathcal{X}$ , and **positive definite**, that is,  $\sum_{i=1}^n \sum_{j=1}^n c_i c_j k(\mathbf{x}_i, \mathbf{x}_j) \ge 0$  for any n > 0, any choice of n objects  $\mathbf{x}_1, \dots, \mathbf{x}_n$  in  $\mathcal{X}$ , and any choice of real numbers  $c_1, \dots, c_n$  in R.



# **Kernel as Inner Product**

The inner product between vectors is the first kernel we encounter.

(called **linear kernel**).

 $\mathcal{X} = R^p \text{ object } \mathbf{x}_i = (x_{i1}, \cdots, x_{ip})^t.$ 

symmetric and positive definite

One is tempted to compare such vectors using their inner product:

for any  $\mathbf{x}_i, \mathbf{x}_j \in \mathbb{R}^p$ ,  $k_L(\mathbf{x}_i, \mathbf{x}_j) := \mathbf{x}_i^T \mathbf{x}_j = \sum_{t=1}^p x_{it} x_{jt}$ .

Represent objects  $\mathbf{x} \in \mathcal{X}$  as a vector  $\phi(\mathbf{x}) \in \mathbb{R}^p$ ,

defining a kernel for any  $\mathbf{x}_i, \mathbf{x}_j \in \mathcal{X}$  by  $k(\mathbf{x}_i, \mathbf{x}_j) = \langle \phi(\mathbf{x}_i), \phi(\mathbf{x}_j) \rangle$ .

**Theorem:** for any kernel k on a space  $\mathcal{X}$ , there exists a Hilbert space  $\mathcal{F}$  and a mapping  $\phi : \mathcal{X} \to \mathcal{F}$  such that  $k(\mathbf{x}_i, \mathbf{x}_j) = \langle \phi(\mathbf{x}_i), \phi(\mathbf{x}_j) \rangle$ , for any  $\mathbf{x}_i, \mathbf{x}_j \in \mathcal{X}$ , where  $\langle u, v \rangle$  represents the dot product in the Hilbert space between any two points  $u, v \in \mathcal{F}$ . (Aronszajn 1950)

Kernels can all be thought of as dot products in feature space  $\mathcal{F}$ . The point  $\mathbf{x} \in \mathcal{X}$  are viewed as point  $\phi(\mathbf{x})$  in  $\mathcal{F}$ .

A Hilbert space is a vector space endowed with a dot product that is complete for the norm induced.R<sup>p</sup> with the classical inner product is an example of a finite-dimensional Hilbert space.



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David Hilbert (01/23/1862 – 02/14/1943)

http://www.hmwu.idv.tw

German mathematician



## Linear kernel and their associated functional space:

Let k be a kernel on a space  $\mathcal{X}$ , to show k is associated with a set of real-valued functions on  $\mathcal{X}$ ,  $\mathcal{H}_k \subset \{f : \mathcal{X} \to R\}$ , endowed with a structure of Hilbert space.

 $\mathcal{X} = R^p$  the functional space is  $f: R^p \to R$  the associated norm is

 $\mathcal{H}_k = \{f(\mathbf{x}) = \mathbf{w}^T \mathbf{x}, \mathbf{w} \in \mathbb{R}^p\} \qquad \qquad \|f\|_{\mathcal{H}_k} = \|\mathbf{w}\| \text{ for } f(\mathbf{x}) = \mathbf{w}^T \mathbf{x}.$ 

The set  $\mathcal{H}_k$  is defined as the set of function  $f : \mathcal{X} \to R$  of the form  $f(\mathbf{x}) = \sum_{i=1}^n \alpha_i k(\mathbf{x}_i, \mathbf{x})$ , for n > 0, a finite number of points  $\mathbf{x}_1, \dots, \mathbf{x}_n \in \mathcal{X}$ , and  $\mathbf{w}$  finite number of weights  $\alpha_1, \dots, \alpha_n \in R$ , together with their limits under the norm  $\| f \|_{\mathcal{H}_k}^2 := \sum_{i=1}^n \sum_{j=1}^n \alpha_i \alpha_j k(\mathbf{x}_i, \mathbf{x}_j)$ .

 $\mathcal{H}_k$  is a Hilbert space, with a dot product defined for two elements  $f(\mathbf{x}) = \sum_{i=1}^n \alpha_i k(\mathbf{x}_i, \mathbf{x})$ and  $g(\mathbf{x}) = \sum_{j=1}^m \alpha'_j k(\mathbf{x}'_j, \mathbf{x})$  by  $\langle f, g \rangle = \sum_{i=1}^n \sum_{j=1}^m \alpha_i \alpha'_j k(\mathbf{x}_i, \mathbf{x}'_j)$ .

The value  $f(\mathbf{x})$  of a function  $f \in \mathcal{H}_k$  at a point  $\mathbf{x} \in \mathcal{X}$  can be expressed

as a dot product in  $\mathcal{H}_k$ ,  $f(\mathbf{x}) = \langle f, k(\mathbf{x}, \cdot) \rangle$ .

taking  $f(\cdot) = k(\mathbf{x}, \cdot)$ : the reproducing property valid for any  $\mathbf{x}_i, \mathbf{x}_j \in \mathcal{X}$ :

$$k(\mathbf{x}_i, \mathbf{x}_j) = \langle k(\mathbf{x}_i, \cdot), k(\mathbf{x}_j, \cdot) \rangle.$$

The functional space  $\mathcal{H}_k$  is usually called the reproducing kernel Hilbert space (RKHS) associated with k.



# **Kernel Trick**

The Hilbert space  $\mathcal{H}_k$  is one possible feature space associated with the kernel k, when we consider the mapping  $\phi : \mathcal{X} \to \mathcal{H}$  defined by  $\phi(\mathbf{x}) := k(\mathbf{x}, \cdot)$ .

- The *kernel Trick* was first published in the 1964 paper *Theoretical foundations* of the potential function method in pattern recognition learning.
- Any algorithm for vectorial data that can be expressed only in terms of *dot products* between vectors can be performed implicitly in the feature space
   associated with any kernel, by replacing each dot product by a kernel evaluation.
- It is a very convenient trick to transform *linear* methods, such as LDA or PCA into *nonlinear* methods, by simply replacing the classic dot product by a more general kernel.
- The kernel trick transforms any algorithm that solely dependents on the dot product between two vectors. Wherever a dot product is used, it is replaced with the kernel function.
- The non-linear algorithm is the linear algorithm operating in the *feature space*.
- *Kernelization*: the operation that transforms a linear algorithm into a more general kernel method.

$$k(x,x') = \langle \Phi(x), \Phi(x') \rangle$$

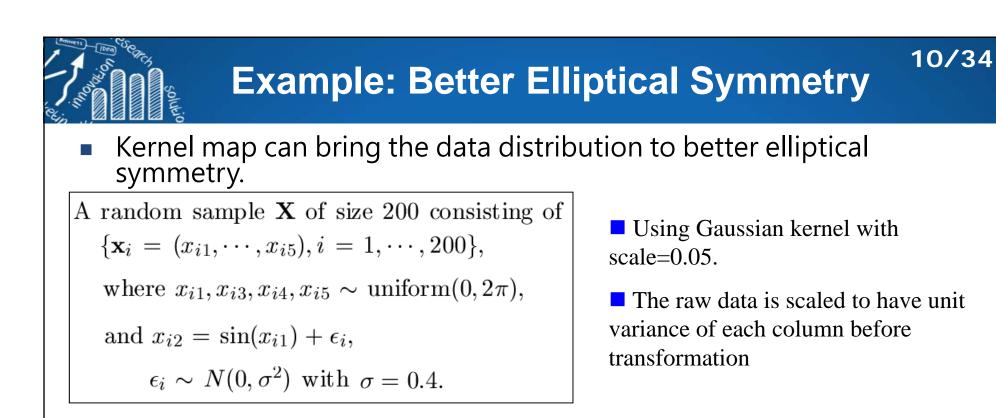
# **Kernel Data: Properties**

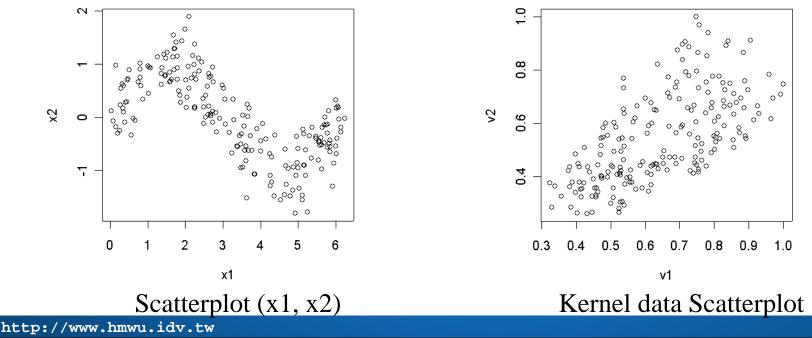
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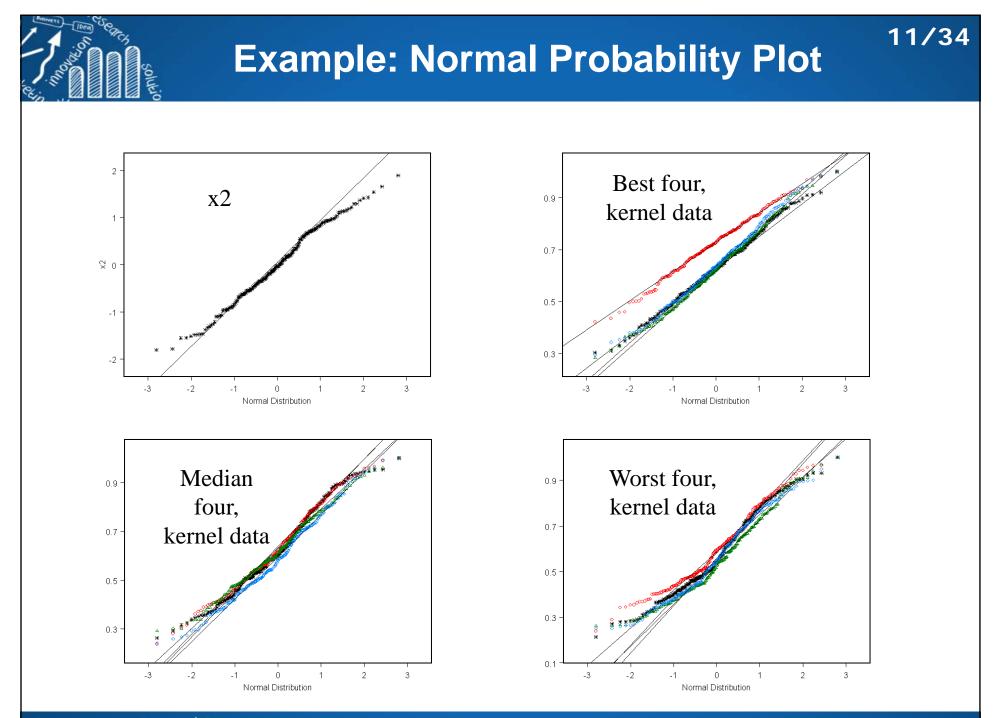
- Raw data on Euclidean space  $R^{p}$ 
  - Kernel data on a RKHS  $H_k$
  - Via a specific statistical notion of classical approach on  $R^{p}$ 
    - Kernel approach on  $H_k$ , which is exactly the classical procedure on kernel data.
  - **Main goal**: Parallel to the classical multivariate statistical analysis, we aim to develop an analysis tool in the Gaussian reproducing kernel Hilbert space.
  - Main advantage: Nonparametric approach with "parametric-plus" computing load.

parametric: classical multivariate analysis procedures.

- plus: kernel data preparation.
- Kernel map can bring the data distribution to better elliptical symmetry. Kernel data are (with empirical and theoretical justification)
  - Better elliptically symmetrically distributed.
  - Better approximately normal (Gaussian)



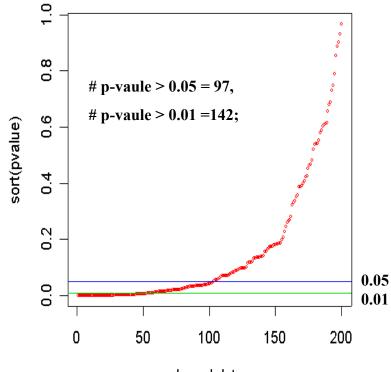




# **Example: Justification of Gaussianity**

**Empirical Justification of Gaussianity:** 

Kolmogorov-Smirnov Test: H<sub>0</sub>: The data follow a normal distribution



kernel data

# **Prepare Your Data to Do the Above Empirical Justification**

## **Theoretical Justification of Gaussianity**

Kernel data  $\{\sqrt{\sigma_n^p} \Gamma_j\}_{j=1}^n$  projected along the random direction h

$$\sqrt{\sigma_n^p}\langle h, \Gamma_1 \rangle_{\mathcal{H}_n}, \cdots, \sqrt{\sigma_n^p}\langle h, \Gamma_n \rangle_{\mathcal{H}_n}.$$

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Let  $\theta_n(h)$  be the empirical distribution of this sequence, assigning probability mass  $n^{-1}$  to each  $\sqrt{\sigma_n^p} \langle h, \Gamma_j \rangle_{\mathcal{H}_n}$ .

**Theorem** Under some conditions, as  $n \to \infty$ , the empirical distribution  $\theta_n(h)$  converges weakly to  $N(0, \tau^2)$  in probability.

### For details:

Huang, S.Y., Hwang, C. R. and Lin, M.H. Kernel Fisher's Discriminant Analysis in Gaussian Reproducing Kernel Hilbert Space.



# **PCA in the Euclidean Space**

# Centered Observations: column vectors $x_i \in \mathbb{R}^N, i = 1, \dots, m$

(Centered meaning:  $\sum_{i=1}^{m} x_i = 0$ )

PCA finds the principal axes by diagonalizing the covariance matrix

$$C = \frac{1}{m} \sum_{j=1}^{m} x_j x_j^\mathsf{T}$$

Note that C is positive definite, and thus can be diagonalized with nonnegative eigenvalues.

$$\lambda \boldsymbol{v} = C \boldsymbol{v}$$

$$C \boldsymbol{v} = rac{1}{m} \sum_{j=1}^m x_j x_j^\mathsf{T} \boldsymbol{v} = \lambda \boldsymbol{v}$$

 $\boldsymbol{v} = \frac{1}{m\lambda} \sum_{j=1}^{m} x_j x_j^{\mathsf{T}} \boldsymbol{v}$  $= \frac{1}{m\lambda} \sum_{j=1}^{m} (x_j \cdot \boldsymbol{v}) x_j$ 

Show that  $(\boldsymbol{x}\boldsymbol{x}^T)\boldsymbol{v} = (\boldsymbol{x}\cdot\boldsymbol{v})\boldsymbol{x}$ 

 $(x_j \cdot \boldsymbol{v})$  is just a scalar

$$\boldsymbol{v} = \sum_{i=1}^{m} \alpha_i x_i$$

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# **Kernel PCA**

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$$\Phi : \mathcal{X} \to \mathcal{H}, \mathbf{x} \mapsto \Phi(\mathbf{x}) \qquad \lambda \sum_{i=1}^{M} \alpha_i (\Phi(\mathbf{x}_k) \cdot \Phi(\mathbf{x}_i)) = \\ \sum_{k=1}^{M} \Phi(\mathbf{x}_k) = 0 \qquad \lambda \sum_{i=1}^{M} \alpha_i (\Phi(\mathbf{x}_k) \cdot \Phi(\mathbf{x}_i)) = \\ \bar{C} = \frac{1}{M} \sum_{j=1}^{M} \Phi(\mathbf{x}_j) \Phi(\mathbf{x}_j)^{\mathsf{T}}, \qquad \frac{1}{M} \sum_{i=1}^{M} \alpha_i (\Phi(\mathbf{x}_k) \cdot \sum_{j=1}^{M} \Phi(\mathbf{x}_j)) (\Phi(\mathbf{x}_j) \cdot \Phi(\mathbf{x}_i)) \\ \lambda V = C V \qquad K_{ij} := (\Phi(\mathbf{x}_i) \cdot \Phi(\mathbf{x}_j)), \\ \lambda (\Phi(\mathbf{x}_k) \cdot \mathbf{V}) = (\Phi(\mathbf{x}_k) \cdot \bar{C} \mathbf{V}) \qquad M \lambda K \alpha = K^2 \alpha, \\ \mathbf{V} = \sum_{i=1}^{M} \alpha_i \Phi(\mathbf{x}_i). \qquad M \lambda \alpha = K \alpha \\ (\mathbf{V}^k \cdot \Phi(\mathbf{x})) = \sum_{i=1}^{M} \alpha_i^k (\Phi(\mathbf{x}_i) \cdot \Phi(\mathbf{x})) \end{cases}$$



# Kernel PCA: kpca {kernlab}

**kernlab**: Kernel-Based Machine Learning Lab

```
> library(kernlab)
> rbf <- rbfdot(sigma = 0.05) #Radial Basis kernel function
> rbf
Gaussian Radial Basis kernel function.
Hyperparameter : sigma = 0.05
> KX <- kernelMatrix(kernel=rbf, x=as.matrix(iris[,1:4])) # calculate kernel matrix
> dim(KX)
[1] 150 150
```

```
test <- sample(1:150, 20)
iris.kpca <- kpca(~., data=iris[-test, -5], kernel="rbfdot", kpar=list(sigma=0.2),</pre>
features=2)
# print the principal component vectors
                                                                                     KPCA for iris data
pcv(iris.kpca)
                                                                        LO.
# plot the data projection on the components
plot(rotated(iris.kpca), col=as.integer(iris[-test, 5]),
                                                                     2nd Principal Component
     xlab="1st Principal Component",
     ylab="2nd Principal Component",
     main="KPCA for iris data")
# embed remaining points
                                                                        ŝ
emb <- predict(iris.kpca, as.matrix(iris[test, -5]))</pre>
points(emb, col=iris[test, 5], pch=17, cex=1.5, asp=1)
                                                                          -10
                                                                                    -5
                                                                                             0
                                                                                    1st Principal Component
```



- Li (1991) introduced the following model
  - $y = f(\beta'_1 \mathbf{x}, \cdots, \beta'_K \mathbf{x}, \epsilon).$

Li, K. C. (1991). Sliced inverse regression for dimensional reduction (with discussion). JASA 86, 316-342.

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y is a univariate variable.

**x** is a random vector with dimension  $p \times 1$ ,  $p \ge K$ .

 $\beta$ 's are vectors with dimension  $p \times 1$ .

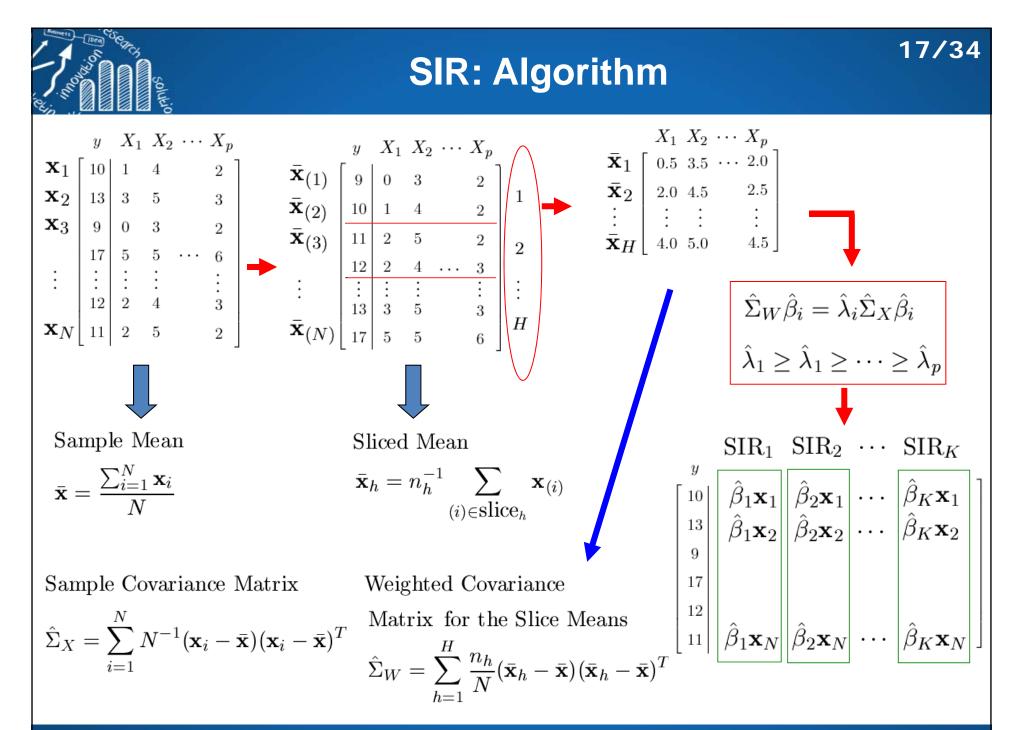
 $\epsilon$  is a random variable independent of  ${\bf x}.$ 

f is an arbitrary function.



- The  $\beta$ 's are referred to effective dimension reduction (*e.d.r.*) or projection directions.

Sliced inverse regression (SIR) is a method for estimating the e.d.r. directions based on y and  $\mathbf{x}$ .





# **SIR: Theorem**

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Linear Design Condition (L.D.C.)

For any b in  $\mathbb{R}^p$ ,

the conditional expectation  $E(b'\mathbf{x}|\beta'_1\mathbf{x},\cdots,\beta'_K\mathbf{x})$  is linear in  $\beta'_1\mathbf{x},\cdots,\beta'_K\mathbf{x}$ ;

 $\blacktriangleright \text{ that is, for some constants } c_0, c_1, \cdots, c_k, \\ E(b'\mathbf{x}|\beta'_1\mathbf{x}, \cdots, \beta'_K\mathbf{x}) = c_0 + c_1\beta'_1\mathbf{x} + \cdots + c_k\beta'_K\mathbf{x}.$ 

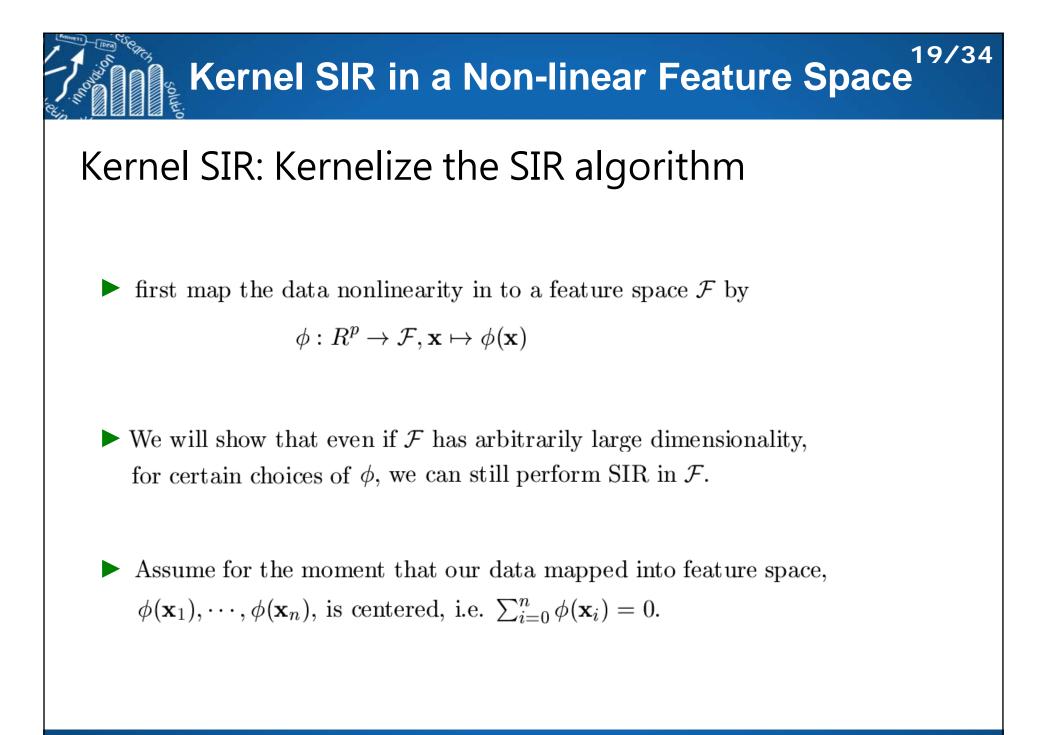
### THEOREM:

under regular conditions, the centered inverse regression curve  $E[\mathbf{x}|y] - E[\mathbf{x}]$ is contained in the linear subspace spanned by  $\beta_k \Sigma_{\mathbf{X}}$   $(k = 1, \dots, K)$ .

## COROLLARY 3.1 (Li, 1991)

Assume that **x** has been standardized to **z**. Then under the model and (3.1), the standardized inverse regression curve  $E(\mathbf{z}|y)$  is contained in the linear space generated by the standardized *e.d.r.* directions  $\theta_1 \ \theta_2 \ \cdots \ \theta_K$ 

The SIR directions  $\mathbf{v_i}$  falls into the *e.d.r* space.



# **KSIR: Algorithm**

We have to find eigenvalues  $\lambda \geq 0$  and eigenvectors  $\boldsymbol{\beta} \in \mathcal{F} \setminus \{0\}$ satisfying  $\Sigma_{wz} \boldsymbol{\beta} = \lambda \Sigma_{zz} \boldsymbol{\beta}$ .

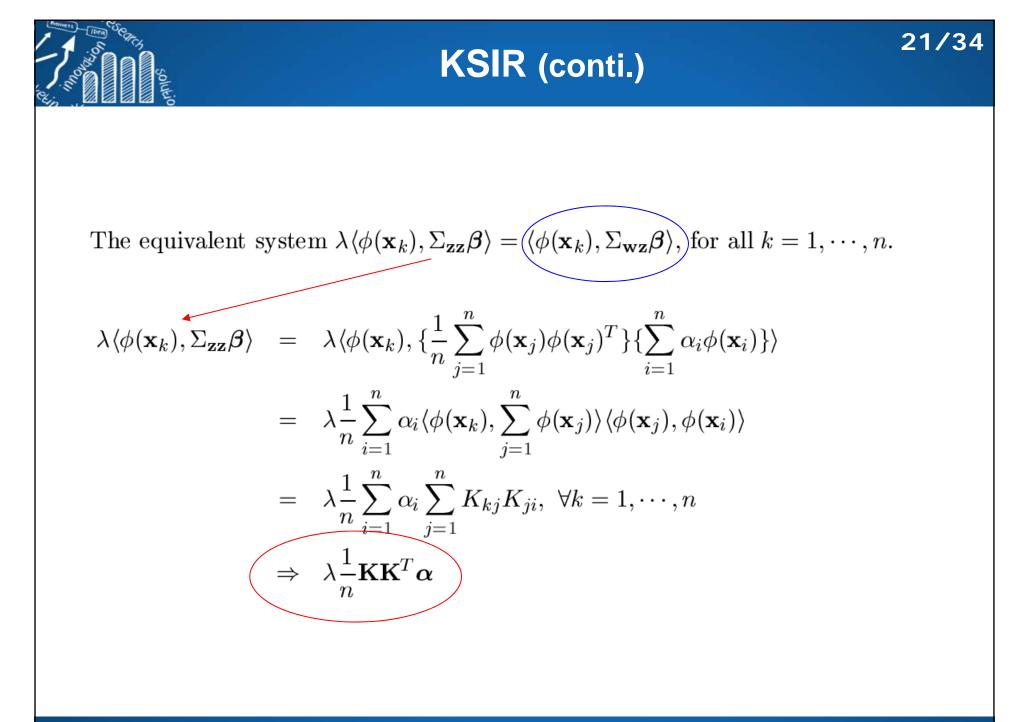
$$\begin{split} \Sigma_{\mathbf{z}\mathbf{z}} &= \frac{1}{n} \sum_{i=1}^{n} \phi(\mathbf{x}_{i}) \phi(\mathbf{x}_{i})^{T}.\\ p_{h} &= \frac{\sum_{i=1}^{n} \delta_{h}(y_{i})}{n} = \frac{n_{h}}{n}, \, \delta_{h}(y_{i}) = 1, \, \text{if} \, y_{i} \in I_{h}, \, \delta_{h}(y_{i}) = 0, \, \text{o.w.} \\ \Sigma_{\mathbf{w}\mathbf{z}} &= \sum_{h=1}^{H} p_{h} \bar{\phi}(\mathbf{m}_{h}) \bar{\phi}(\mathbf{m}_{h})^{T}.\\ \bar{\phi}(\mathbf{m}_{h}) &= \frac{1}{np_{h}} \sum_{i=1}^{n} \phi(\mathbf{x}_{i}) \delta_{h}(y_{i}) \end{split}$$

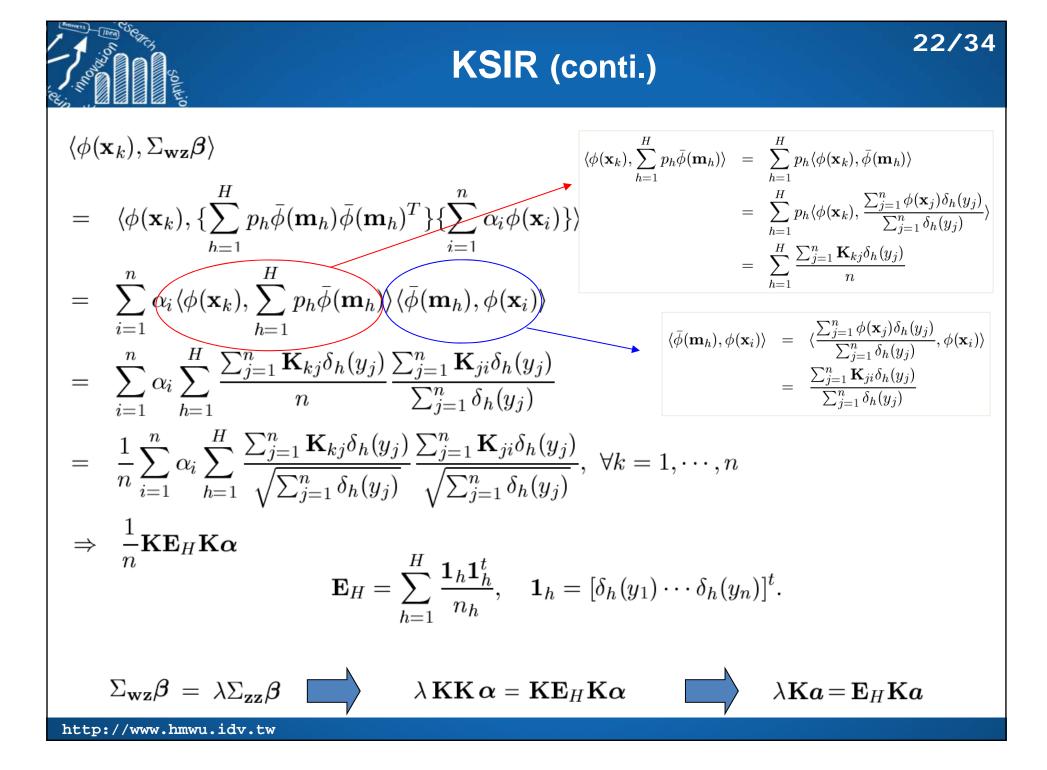
All solutions  $\boldsymbol{\beta}$  lie in span  $\{\phi(\mathbf{x}_1), \cdots, \phi(\mathbf{x}_n)\}.$ 

The equivalent system  $\lambda \langle \phi(\mathbf{x}_k), \Sigma_{\mathbf{z}\mathbf{z}} \boldsymbol{\beta} \rangle = \langle \phi(\mathbf{x}_k), \Sigma_{\mathbf{w}\mathbf{z}} \boldsymbol{\beta} \rangle$ , for all  $k = 1, \dots, n$ .

▶ there exists  $\alpha_1, \dots, \alpha_n$  such that  $\beta = \sum_{i=1}^n \alpha_i \phi(\mathbf{x}_i)$ .

Define 
$$\mathbf{K} := \{\mathbf{k}_{ij} = \langle \phi(\mathbf{x}_i), \phi(\mathbf{x}_j) \rangle \}_{n \times n}.$$





# **Normalization and Projection**

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Let  $\lambda_1 \geq \cdots \geq \lambda_n$  denote the eigenvalues, and  $\boldsymbol{\alpha}_1, \cdots, \boldsymbol{\alpha}_n$ the corresponding complete set of eigenvectors, with  $\lambda_t$ being the first nonzero eigenvalues.

We normalize  $\boldsymbol{\alpha}_1, \dots, \boldsymbol{\alpha}_n$  by requiring that the corresponding vectors in  $\mathcal{F}$  be normalized:  $\langle \boldsymbol{\beta}_k, \boldsymbol{\beta}_k \rangle = 1$  for all  $k = 1, \dots, t$ .

Normalization Condition:

$$1 = \sum_{i=1}^{n} \sum_{j=1}^{n} \alpha_i^k \alpha_j^k \langle \phi(\mathbf{x}_i), \phi(\mathbf{x}_j) \rangle = \langle \boldsymbol{\alpha}^k, \mathbf{K} \boldsymbol{\alpha}^k \rangle = \lambda_k \langle \boldsymbol{\alpha}^k, \boldsymbol{\alpha}^k \rangle$$

**Projections on the eigenvectors**  $\beta_k$  in  $\mathcal{F}$ ,  $k = 1, \dots, t$ :

Let **x** be a test point, with an image  $\phi(\mathbf{x})$  in  $\mathcal{F}$ , then

$$\langle \boldsymbol{\beta}_k, \phi(\mathbf{x}) \rangle = \sum_{i=1}^n \alpha_i^k \langle \phi(\mathbf{x}_i), \phi(\mathbf{x}) \rangle = \sum_{i=1}^n \alpha_i^k \mathbf{K}(\mathbf{x}_i, \mathbf{x})$$



# **Centering in Feature Space**

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The mapped data is centered in  $\mathcal{F}$ ,  $\sum_{i=1}^{n} \phi(\mathbf{x}_i) = 0$ .

- The points  $\tilde{\phi}(\mathbf{x}_i) := \phi(\mathbf{x}_i) \frac{1}{n} \sum_{i=1}^n \phi(\mathbf{x}_i)$  will be centered.
- Define  $\tilde{\mathbf{K}} := \langle \tilde{\phi}(\mathbf{x}_i), \tilde{\phi}(\mathbf{x}_i) \rangle$  in  $\mathcal{F}$ .

$$\tilde{\mathbf{K}} = \mathbf{K} - I_n \mathbf{K} - \mathbf{K} I_n + I_n \mathbf{K} I_n, \ (I_n)_{ij} = 1/n.$$

## **For Training Data**

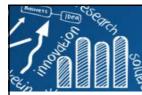
 $K_{tr} \leftarrow \operatorname{kernelMatrix}(poly, X_{tr})$ 

$$K_{tr.c} \leftarrow K_{tr} - \mathbf{1}_{tr} K_{tr} - K_{tr} \mathbf{1}_{tr} + \mathbf{1}_{tr} K_{tr} \mathbf{1}_{tr}$$

## **For Testing Data**

$$K_{te} \leftarrow \operatorname{kernelMatrix}(poly, \boldsymbol{X}_{te}, \boldsymbol{X}_{tr})$$

$$K_{te.c} \leftarrow K_{te} - \mathbf{1}_{te} K_{tr} - K_{te} \mathbf{1}_{tr} + \mathbf{1}_{te} K_{tr} \mathbf{1}_{tr}$$



## **Reduced Features**

- we are not working in the **full feature space**, but just in a comparably small linear subspace of it, whose dimension equals at most the number of observations.
- Working in a space whose dimension equals the number of observations can pose difficulties.
- To deal with these, one can either use only a subset of the extracted features, or use some other form of capacity control or regularization.

Data Transform Dialog	
Data Set	New Data
SuYun-sinData.txt 🚽 [ 200 x 5 ]	tSuYun-sinData.txt [ 200 x 200 ]
Transform Methods	
Kernel Transform 👻	
Settings	
🖌 Standardize Data by Columns	
Kernel Type with Parameters	
Gaussian RBF 👻 degree: 2	scale: 0.05 offset: 0
Sampling Columns	
○ Random 200	
	OK Cancel

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#### For Theoretical details:

Lee, Y.J. and Huang, S.Y. (2006), Reduced support vector machines: a statistical theory, *IEEE Transactions on Neural Networks*, accepted. http://dmlab1.csie.ntust.edu.tw/downloads

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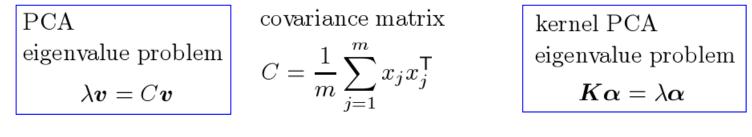
# **Relations Towards Other Methods**

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## SIR vs. KSIR

- KSIR generalizes SIR to a nonlinear one by kernelization of the SIR algorithm.
- It finds nonlinear d.r. subspace, a central d.r. subspace in  $H_k$
- A semiparametric method.
- SIR: spectrum analysis of cov(E[x|y]) wrt cov(x)
- **KSIR**: spectrum analysis of a generalized association measure.

## KSIR vs. KPCA





KSIR

PCA performed on the random vector  $E(\mathbf{x}|y)$  instead of  $\mathbf{x}$ . PCA performed on the random vector  $E(\phi(\mathbf{x})|y)$  instead of  $\phi(\mathbf{x})$ .

# **Relations Towards Other Methods**

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## KSIR vs. KFDA

$$\max_{a} \frac{a^{t} \Sigma_{B} a}{a^{t} \Sigma_{W} a} \implies \Sigma_{B} a = \gamma_{i} \Sigma_{W} a, \quad \gamma_{1} \ge \gamma_{2} \ge \cdots \ge \gamma_{p}$$
$$\implies \Sigma_{\mathbf{x}\mathbf{x}} = \Sigma_{B} + \Sigma_{W} \implies \Sigma_{B} a_{i} = \frac{\gamma_{i}}{1 + \gamma_{i}} \Sigma_{\mathbf{x}\mathbf{x}} a_{i}$$
$$\Sigma_{\mathbf{w}\mathbf{x}} \beta_{j} = \lambda_{j} \Sigma_{\mathbf{x}\mathbf{x}} \beta_{j}$$
$$\implies \lambda_{i} = \gamma/(1 + \gamma) \text{ and } a_{i} \propto \beta_{i},$$
Chen, C. H., and Li, K. C. (2001)

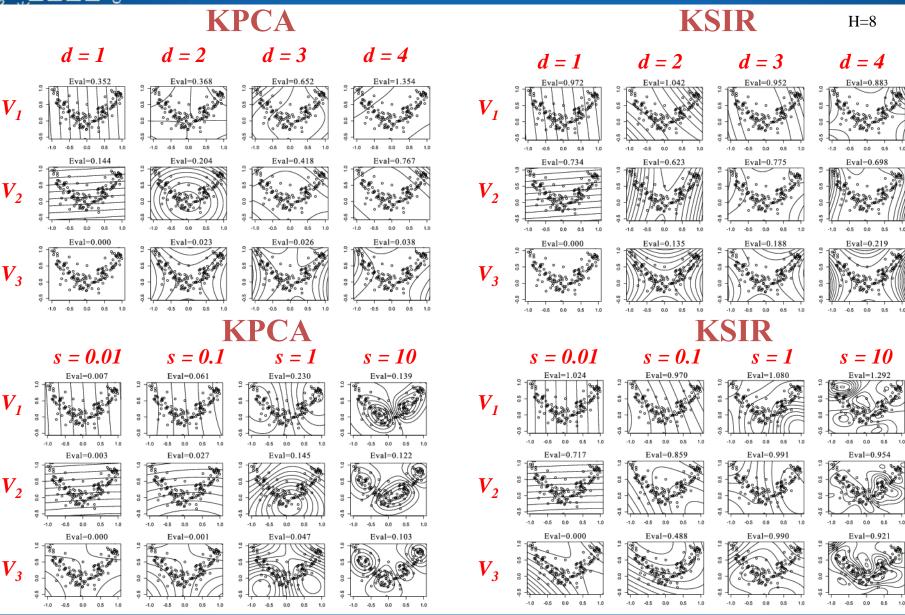
### KSIR vs. KCCA

Kernel Fisher discriminant Analysis as special case of CCA. (Kuss, M. and Graepel, T: The Geometry Of Kernel Canonical Correlation Analysis. (108), Max Planck Institute for Biological Cybernetics, Tübingen, Germany (May 2003)

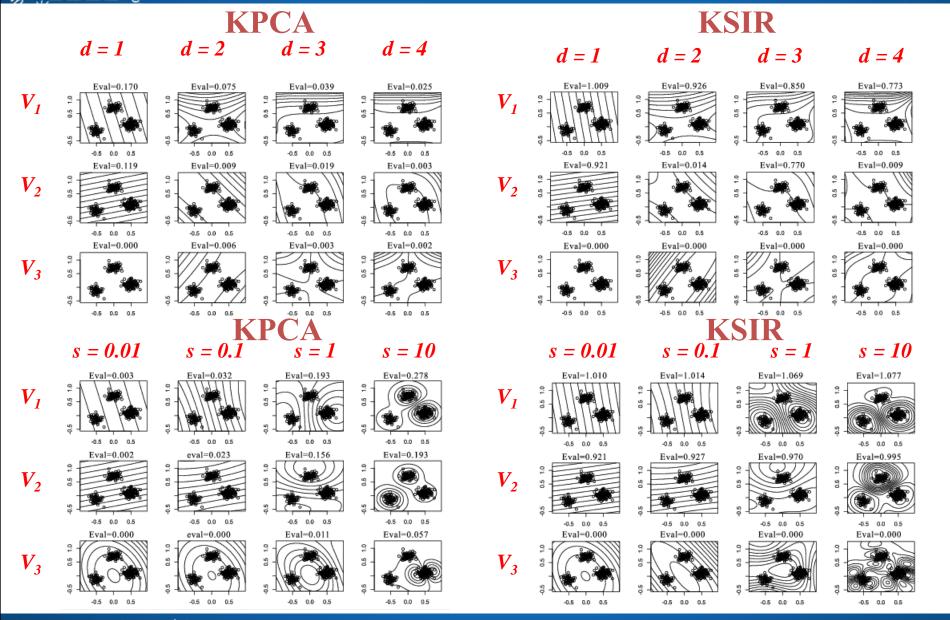


# **Visualization: Square Data (150x2)**

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Visualization: Three Clusters Data (220x2)

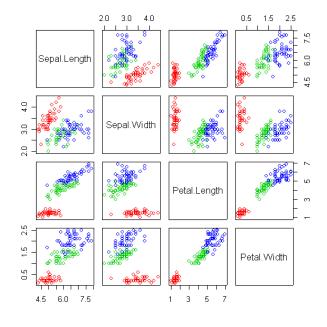


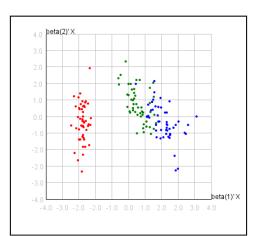
# Visualization: Iris Data (150x4)

beta(2)')

■ The sepal length, sepal width, petal length, and petal width are measured in centimeters on 50 iris specimens from each of three species, *Iris setosa*, *I. versicolor*, and *I. virginica*. Fisher (1936)

Solute

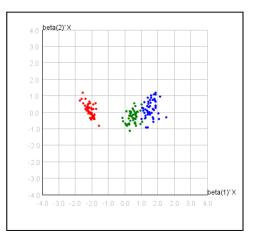




**PCA** 

## SIR

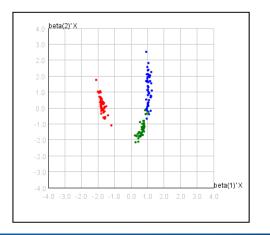
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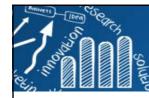


Gaussian s=0.05

eta(1)'X

## **KSIR**





# Visualization: Wine Data (178x18)

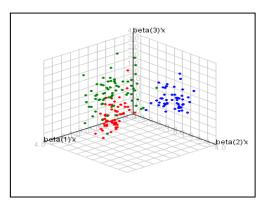
## PCA

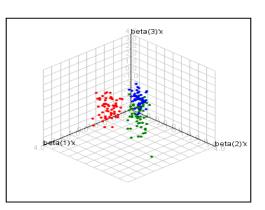
## SIR

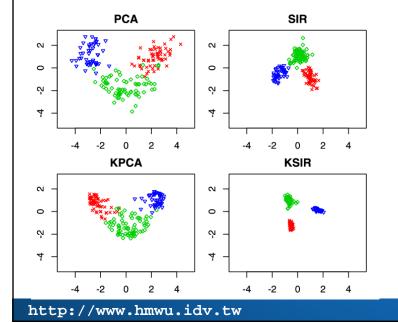
Wine data (n=178) are the results of a chemical analysis of wines grown in the same region in Italy but derived from three different cultivars.

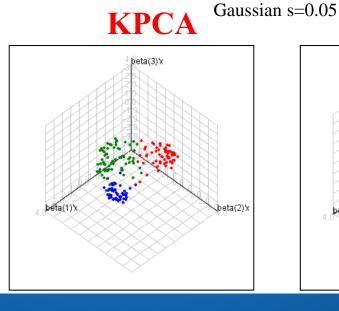
The analysis determined the quantities of 13 constituents found in each of the three types of wines.

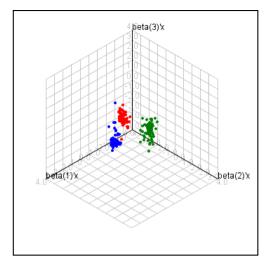
Past Usage
 RDA : 100%, QDA 99.4%,
 LDA 98.9%, 1NN 96.1%
 (z-transformed data, loo)











**KSIR** 

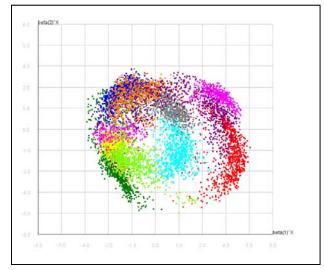
# Visualization: Pendigit Data (7494x16)

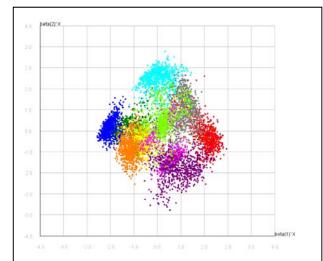
PCA

SIR

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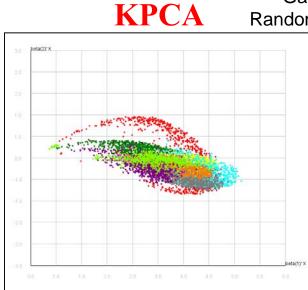
- Pen-based recognition of handwritten Digits
- 7494 instances, 16 attributes
- 10 classes

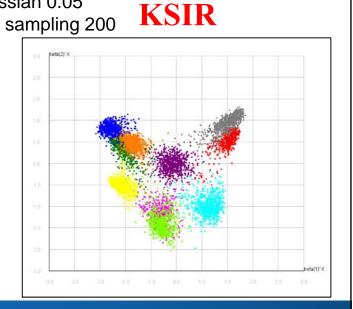




#### Gaussian 0.05 Random sampling 200

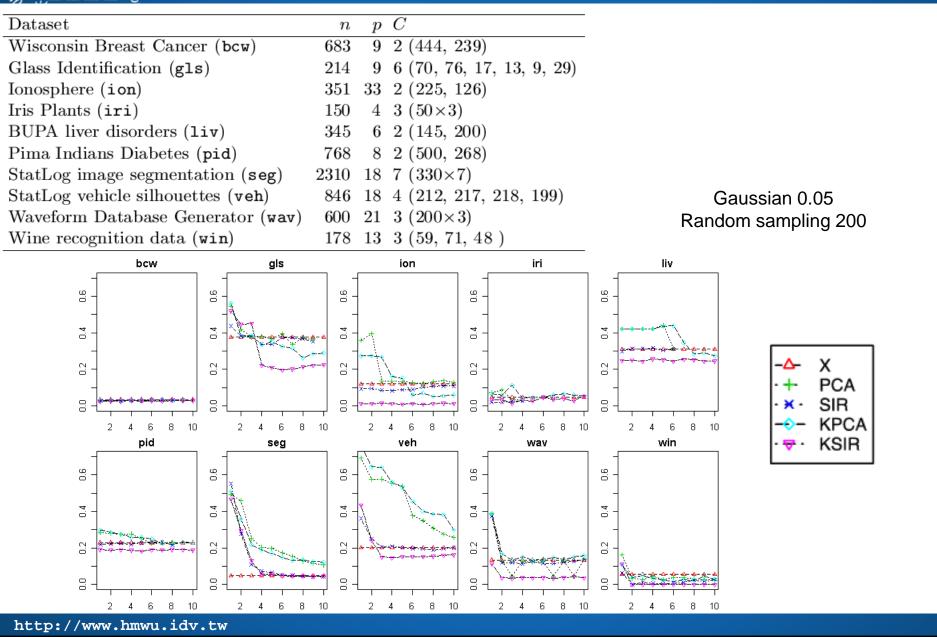








# **Classification: UCI Data Sets**





# **Classification: Microarray Data Sets**

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#### Brain Colon Leukemia Dataset Publication np72 3571 Golub *et al.*(1999) Leukemia 0.0 R 0.30 Alon *et al.*(1999) Colon 62 2000 0 Singh et al.(2002)102 6033 Prostate Lymphoma Alizadeh et al. (2000) 62 4026 0.20 0.20 SRBCT Khan et al. (2001) 6323080.4 Pomeroy et al. (2002) Brain 42 5597 0.10 0.10 0.2 Dataset CResponse 80 8.0 0.0 Subtypes of leukemia Leukemia 2(47, 25)2(22, 40)Tumor/normal tissue Colon 6 10 2 6 8 10 8 10 2 4 8 4 2 6 4 Lymphoma Prostate SRBCT Tumor/normal tissue Prostate 2(50, 52)Subtypes of lymphoma Lymphoma 3(42, 9, 11)۵ Ö 8.0 Different tumor types SRBCT 4 (23, 20, 12, 8) 0.4 чO ö Brain 5(10, 10, 10, 4, 8)Different tumor types 0.4 0.3 0.20 0.3 0.2 Х -A-0.2 0.10 KPCA.s0.05 <u>.</u> KPCA.d3 --<del>X</del>--0.1 KSIR.s0.05 ↔ 8.0 0.0 0 Ö KSIR.d3 -8-2 6 8 10 2 6 8 10 2 6 8 10 4 4 Δ