

Nonlinear model

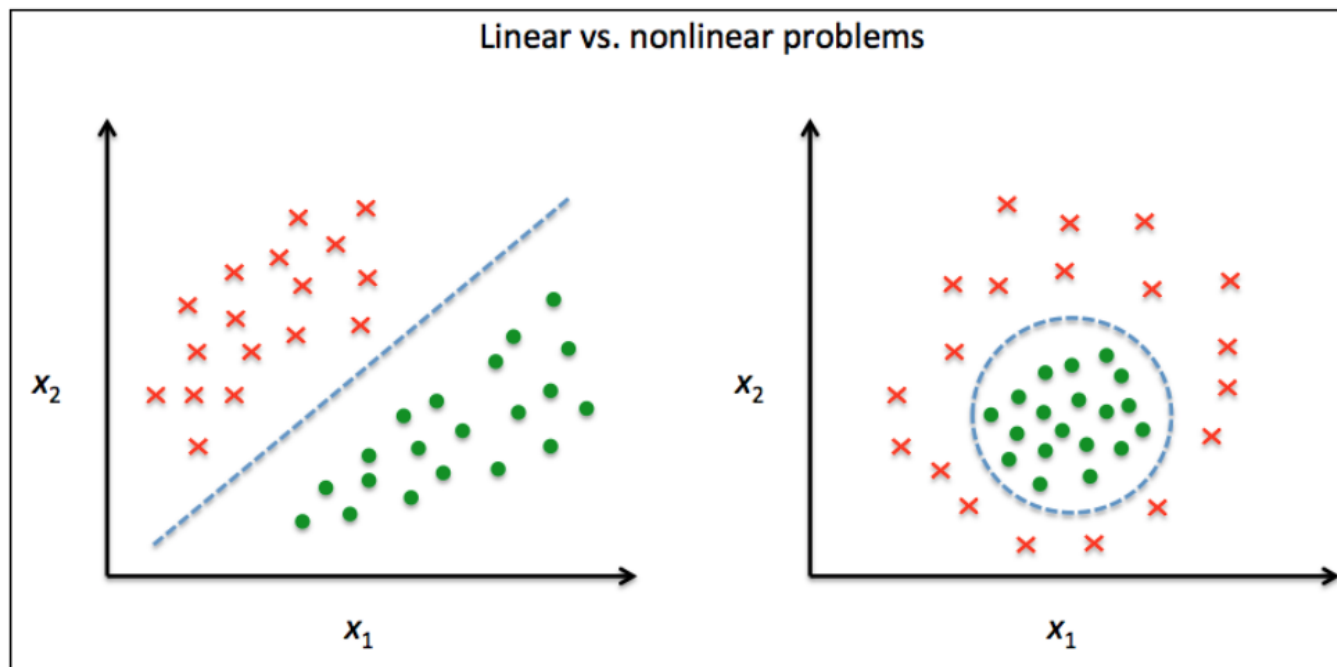
via feature transform & kernel trick

DeepNK Machine Learning Webinar

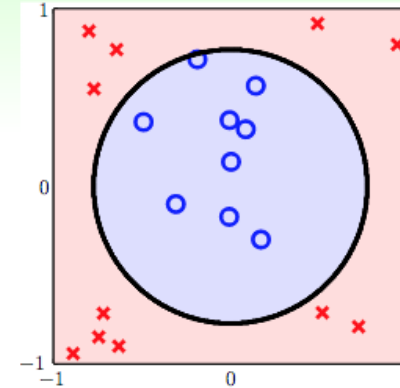
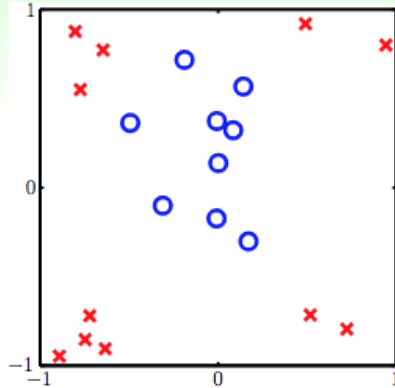
(<https://hackmd.io/CwBgTAxgbARgZgQwLQEYDMYCSTRqkgTgFM584DoB2IIMADgBMI6g>)

Linear model is limited ...

- On some dataset, the linear model just performed poorly during training phase
- Non-trivially linear separable data
- Essentially training data are non-linear



Circular Separable



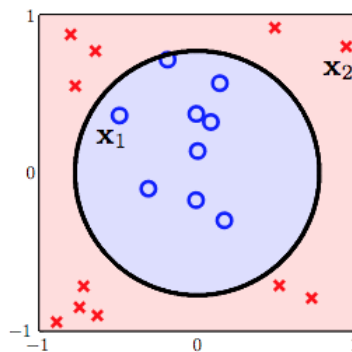
- \mathcal{D} not linear separable
- but **circular separable** by a circle of radius $\sqrt{0.6}$ centered at origin:

$$h_{\text{SEP}}(\mathbf{x}) = \text{sign} \left(-x_1^2 - x_2^2 + 0.6 \right)$$

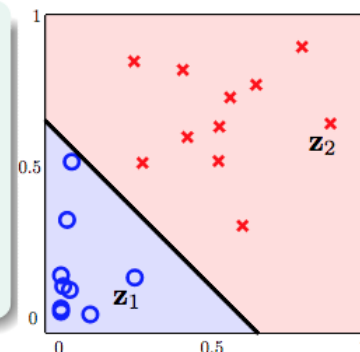
Often we want to **capture nonlinear patterns** ...

Circular Separable and Linear Separable

$$\begin{aligned}
 h(\mathbf{x}) &= \text{sign} \left(\underbrace{0.6}_{\tilde{w}_0} \cdot \underbrace{1}_{z_0} + \underbrace{(-1)}_{\tilde{w}_1} \cdot \underbrace{x_1^2}_{z_1} + \underbrace{(-1)}_{\tilde{w}_2} \cdot \underbrace{x_2^2}_{z_2} \right) \\
 &= \text{sign} \left(\tilde{\mathbf{w}}^T \mathbf{z} \right)
 \end{aligned}$$



- $\{(\mathbf{x}_n, y_n)\}$ circular separable
 $\implies \{(\mathbf{z}_n, y_n)\}$ linear separable
- $\mathbf{x} \in \mathcal{X} \xrightarrow{\Phi} \mathbf{z} \in \mathcal{Z}$:
(nonlinear) feature transform Φ



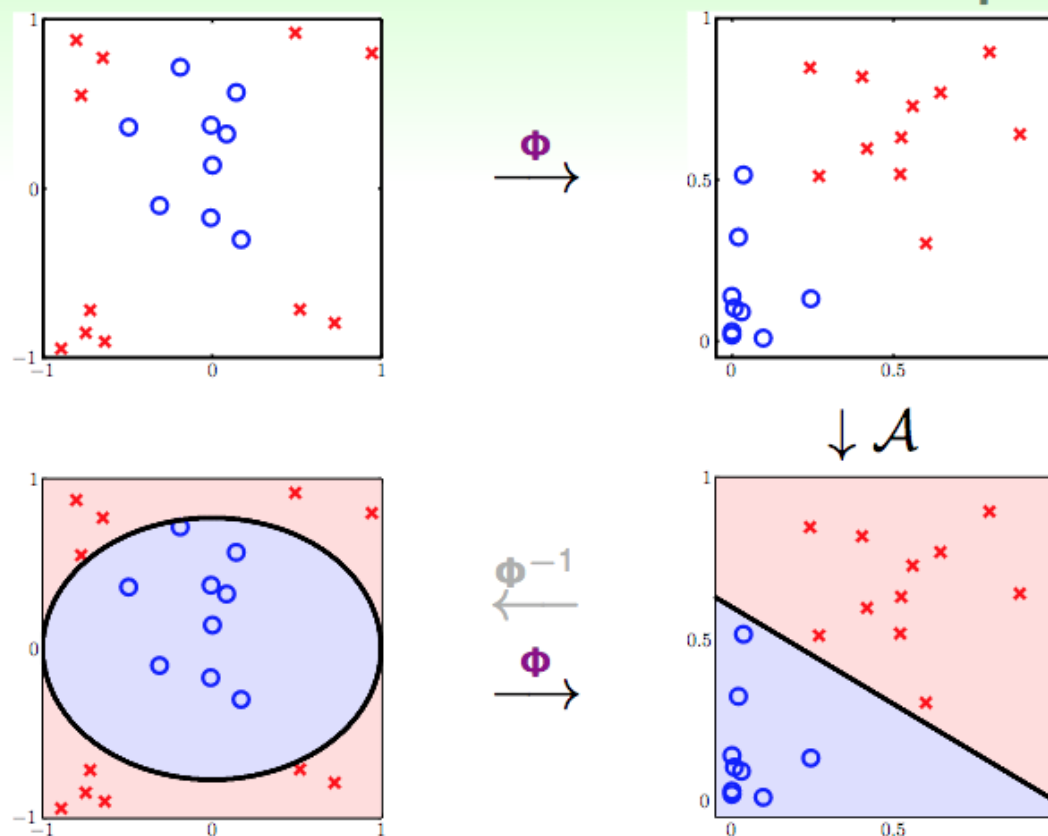
From: Hsuan-Tien Lin's ML Foundation (<https://www.youtube.com/watch?v=8pQ06pku1xA>)

Idea of nonlinear transform

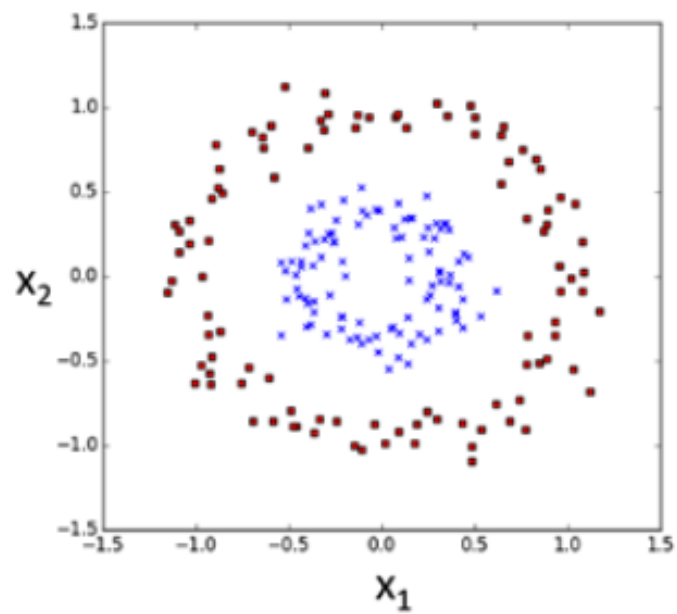
Instead of direct linear weighting, we can project/map features by non-linear feature transform $\Phi : \mathcal{X} \rightarrow \mathcal{F}$ and then apply linear weighting to these mapped features.

- So we can apply good-old linear classification algorithms (e.g. Perceptron, Logistic regression, etc ...)
- An inverse mapping is not required, but we human usually define bijection for good.

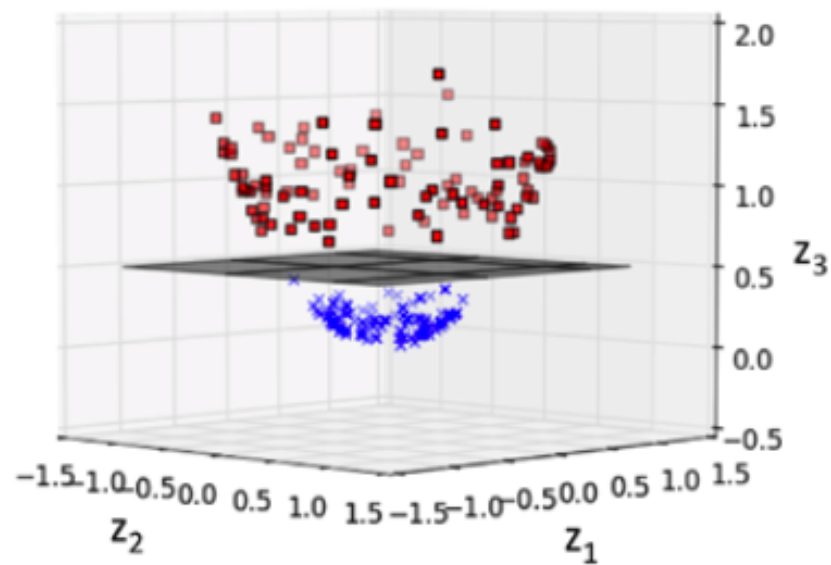
The Nonlinear Transform Steps



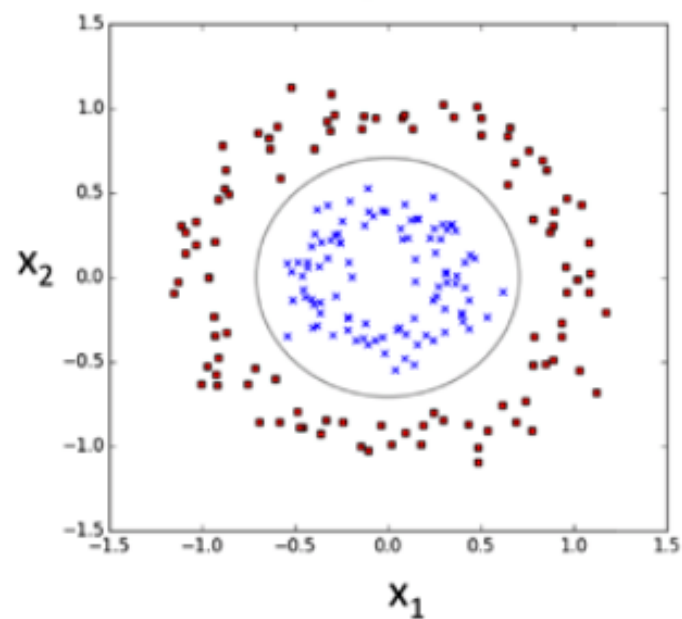
- 1 transform original data $\{(\mathbf{x}_n, y_n)\}$ to $\{(\mathbf{z}_n = \Phi(\mathbf{x}_n), y_n)\}$ by Φ
- 2 get a good perceptron $\tilde{\mathbf{w}}$ using $\{(\mathbf{z}_n, y_n)\}$ and your favorite linear classification algorithm \mathcal{A}
- 3 return $g(\mathbf{x}) = \text{sign}(\tilde{\mathbf{w}}^T \Phi_2(\mathbf{x}))$



ϕ



ϕ^{-1}



Nonlinear and higher dimensionality

The transform does not need to increase the dimension of feature vector space (As written in the book). But one may argue that:

A circle in 2D is one kind of conic section in 3D. So it is in a sense that a nonlinear transform projects feature vectors into a higher dimensional space.

And usually, a linear separator in higher dimension implies nonlinear separator in the original space.

But, No free lunch ...

As we know that nonlinear model increase the expressive power to describe the data, but it comes with price ...

- The price of **computation/storage**
 - But **kernel tricks** will help, see later !!
- The price of **model complexity**
 - The buzzword: **overfitting**

Computation/Storage Price

Q -th order polynomial transform: $\Phi_Q(\mathbf{x}) = \left(\begin{array}{l} 1, \\ x_1, x_2, \dots, x_d, \\ x_1^2, x_1 x_2, \dots, x_d^2, \\ \dots, \\ x_1^Q, x_1^{Q-1} x_2, \dots, x_d^Q \end{array} \right)$

$\underbrace{1}_{\tilde{w}_0} + \underbrace{\tilde{d}}_{\text{others}}$ dimensions
= # ways of $\leq Q$ -combination from d kinds with repetitions
= $\binom{Q+d}{Q} = \binom{Q+d}{d} = O(Q^d)$
= efforts needed for computing/storing $\mathbf{z} = \Phi_Q(\mathbf{x})$ and $\tilde{\mathbf{w}}$

Q large \implies **difficult to compute/store**

Idea of Kernel method (tricks)

We know from above that such a **nonlinear transform** is usually costly to compute and store, especially in much higher dimensions.

In some learning algorithms (https://en.wikipedia.org/wiki/Instance-based_learning), they may compute the **dot product** or **similarity measure** between data samples.

Is there any way to do **nonlinear transform** and **similarity measure** “at the same time”?

Def. of kernel function

Let $\mathbf{x} = [x_1, x_2, \dots, x_m]^T$ be the sample vector in m -dimension input space. And the nonlinear transform $\Phi : \mathcal{X} \rightarrow \mathcal{F}$ maps $\mathbf{x} \in \mathcal{X}$ into $\phi(\mathbf{x}) \in \mathcal{F}$

The kernel function $\kappa : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R}$

$$k(\mathbf{x}, \mathbf{x}') = \phi(\mathbf{x})^T \phi(\mathbf{x}')$$

computes **dot product similarity** in the transformed feature space \mathcal{Z} given two sample vectors \mathbf{x} and \mathbf{x}' in original input space

Properties of kernel function

- $k(\mathbf{x}, \mathbf{x}')$ must be a proper inner product in feature space \mathcal{Z} . So \mathcal{Z} is an inner product space (https://en.wikipedia.org/wiki/Inner_product_space).
 - If \mathcal{Z} is a complete vector space then it is called Hilbert space (https://en.wikipedia.org/wiki/Hilbert_space)
- Can just *any* function be used as a kernel function ?
 - No, It must satisfy **Mercer's Condition** (https://en.wikipedia.org/wiki/Mercer%27s_condition)
- The kernel will do **implicit** feature transform

Mercer's Condition

- For k to be a kernel function
 - There must exist a Hilbert Space \mathcal{F} for which k defines a dot product
 - The above is true if K is a **positive definite function**

$$\int d\mathbf{x} \int d\mathbf{z} f(\mathbf{x}) k(\mathbf{x}, \mathbf{z}) f(\mathbf{z}) > 0 \quad (\forall f \in L_2)$$

- This is Mercer's Condition
- Let k_1, k_2 be two kernel functions then the following are as well:
 - $k(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z}) + k_2(\mathbf{x}, \mathbf{z})$: direct sum
 - $k(\mathbf{x}, \mathbf{z}) = \alpha k_1(\mathbf{x}, \mathbf{z})$: scalar product
 - $k(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z}) k_2(\mathbf{x}, \mathbf{z})$: direct product
 - Kernels can also be constructed by composing these rules

Kernels as High Dimensional Feature Mapping

- Consider two examples $\mathbf{x} = \{x_1, x_2\}$ and $\mathbf{z} = \{z_1, z_2\}$
- Let's assume we are given a function k (kernel) that takes as inputs \mathbf{x} and \mathbf{z}

$$\begin{aligned}k(\mathbf{x}, \mathbf{z}) &= (\mathbf{x}^\top \mathbf{z})^2 \\&= (x_1 z_1 + x_2 z_2)^2 \\&= x_1^2 z_1^2 + x_2^2 z_2^2 + 2x_1 x_2 z_1 z_2 \\&= (x_1^2, \sqrt{2}x_1 x_2, x_2^2)^\top (z_1^2, \sqrt{2}z_1 z_2, z_2^2) \\&= \phi(\mathbf{x})^\top \phi(\mathbf{z})\end{aligned}$$

- The above k **implicitly** defines a mapping ϕ to a higher dimensional space

$$\phi(\mathbf{x}) = \{x_1^2, \sqrt{2}x_1 x_2, x_2^2\}$$

- Note that we didn't have to define/compute this mapping
- Simply defining the kernel a certain way gives a higher dim. mapping ϕ
- Moreover the kernel $k(\mathbf{x}, \mathbf{z})$ also computes the dot product $\phi(\mathbf{x})^\top \phi(\mathbf{z})$
 - $\phi(\mathbf{x})^\top \phi(\mathbf{z})$ would otherwise be much more expensive to compute explicitly
- All kernel functions have these properties

Kernel PCA

Simply put: Apply PCA on the projected features $\phi(\mathbf{x})$, but use **kernel tricks** to avoid the explicit Φ transform. Before we add the kernel tricks, let's recap *standard* PCA:

- Standardize the dataset
- Compute the covariance matrix Σ on the dataset
- Find the eigenvalues and eigenvectors of Σ
- Select m-th principal components
- Project the dataset by these PCs

Projected covariance matrix

After standardization, each element σ_{jk} of $m \times m$ covariance matrix Σ is

$$\sigma_{jk} = \frac{1}{n} \sum_{i=1}^n \mathbf{x}_j^{(i)} \mathbf{x}_k^{(i)}$$

Where $\mathbf{x}^{(i)} = [x_1^{(i)}, x_2^{(i)}, \dots, x_m^{(i)}]^T$ represents a mean-centered sample vector with m features. Reader may quickly find that Σ is also the sum of n $m \times m$ matrices of $\mathbf{x}^{(i)} \mathbf{x}^{(i)T}$ (outer product of $\mathbf{x}^{(i)}$ to itself)

So we can write the covariance matrix in this form:

$$\begin{aligned}\Sigma &= \frac{1}{n} \sum_{i=1}^n \mathbf{x}^{(i)} \mathbf{x}^{(i)T} = \frac{1}{n} [\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(n)}] \begin{bmatrix} \mathbf{x}^{(1)T} \\ \vdots \\ \mathbf{x}^{(n)T} \end{bmatrix} \\ &= \frac{1}{n} X^T X, \quad \mathbf{x}_j^{(i)} \in X_{n \times m}, \quad i \leq n, \quad j \leq m\end{aligned}$$

Replace each $\mathbf{x}^{(i)}$ into projected $\phi(\mathbf{x}^{(i)})$ for us to apply PCA on the projected space:

$$\begin{aligned}\Sigma_{\Phi} &= \frac{1}{n} \sum_{i=1}^n \phi(\mathbf{x}^{(i)}) \phi(\mathbf{x}^{(i)})^T \\ &= \frac{1}{n} [\phi(\mathbf{x}^{(1)}), \dots, \phi(\mathbf{x}^{(n)})] \begin{bmatrix} \phi(\mathbf{x}^{(1)})^T \\ \vdots \\ \phi(\mathbf{x}^{(n)})^T \end{bmatrix} \\ &= \frac{1}{n} \phi(X)^T \phi(X)\end{aligned}$$

The kernel (Gram) matrix

We know that directly compute the projected covariance matrix $\Sigma_{\Phi} = \frac{1}{n}\phi(X)^T\phi(X)$ is costly or sometimes impossible. Time for the **kernel tricks**!

We can use the kernel function $k(\mathbf{x}, \mathbf{x}') = \phi(\mathbf{x})^T\phi(\mathbf{x}')$ to construct the $n \times n$ **kernel matrix** K such that each element $K_{ij} = \phi(\mathbf{x}^{(i)})^T\phi(\mathbf{x}^{(j)})$.

How we relate K to the projected PCA:

$$\Sigma_{\Phi} \mathbf{v} = \lambda \mathbf{v} ?$$

We can rewrite the kernel matrix K as:

$$K = \begin{bmatrix} \phi(\mathbf{x}^{(1)})^T \\ \vdots \\ \phi(\mathbf{x}^{(n)})^T \end{bmatrix} [\phi(\mathbf{x}^{(1)}), \dots, \phi(\mathbf{x}^{(n)})] = \phi(X)\phi(X)^T$$

- $K = \phi(X)\phi(X)^T$ ($n \times n$ matrix)
 $\neq \phi(X)^T\phi(X)$ ($? \times ?$ matrix)
- K is a positive semi-definite matrix by the symmetry property of dot product.
And we can solve the eigenproblem of it: $K\mathbf{u} = \lambda\mathbf{u}$.

Multiply $\phi(X)^T$ to the both side of $K\mathbf{u} = \lambda\mathbf{u}$:

$$\begin{aligned}\phi(X)^T K\mathbf{u} &= \phi(X)^T [\phi(X)\phi(X)^T]\mathbf{u} = \lambda\phi(X)^T\mathbf{u} \\ \implies [\phi(X)^T\phi(X)]\phi(X)^T\mathbf{u} &= \lambda\phi(X)^T\mathbf{u}\end{aligned}$$

- This means $\phi(X)^T\mathbf{u}$ is the eigenvector of $\phi(X)^T\phi(X) = n\Sigma_\Phi$.

Voila! We just relate K to the projected PCA

- However, the norm of $\phi(X)^T\mathbf{u}$ may not be 1 (not ortho"normal")

Projecting test samples

The eigenvector \mathbf{v} of $\phi(X)^T\phi(X)$ can be computed by the eigenvalue λ of K :

$$\begin{aligned}\mathbf{v} &= \frac{1}{\|\phi(X)^T\mathbf{u}\|}\phi(X)^T\mathbf{u} = \frac{1}{\sqrt{\mathbf{u}^T\phi(X)\phi(X)^T\mathbf{u}}}\phi(X)^T\mathbf{u} \\ &= \frac{1}{\sqrt{\mathbf{u}^T(\lambda\mathbf{u})}}\phi(X)^T\mathbf{u} = \frac{1}{\sqrt{\lambda}}\phi(X)^T\mathbf{u}\end{aligned}$$

Uh... but we don't know $\phi(X)^T$... Should we compute \mathbf{v} for transforming test sample \mathbf{x}' to the principal components of feature space?

No, we use **kernel tricks** again!

The principal component projection (dot product) of the test sample $\phi(\mathbf{x}')$ in feature space is:

$$\begin{aligned} \mathbf{v}^T \phi(\mathbf{x}') &= \left[\frac{1}{\sqrt{\lambda}} \phi(X)^T \mathbf{u} \right]^T \phi(\mathbf{x}') = \frac{1}{\sqrt{\lambda}} \mathbf{u}^T \phi(X) \phi(\mathbf{x}') \\ &= \frac{1}{\sqrt{\lambda}} \mathbf{u}^T \begin{bmatrix} \phi(\mathbf{x}^{(1)})^T \\ \vdots \\ \phi(\mathbf{x}^{(n)})^T \end{bmatrix} \phi(\mathbf{x}') = \frac{1}{\sqrt{\lambda}} \mathbf{u}^T \begin{bmatrix} k(\mathbf{x}^{(1)}, \mathbf{x}') \\ \vdots \\ k(\mathbf{x}^{(n)}, \mathbf{x}') \end{bmatrix} \end{aligned}$$

It needs original training dataset to project new samples, a kind of instance-based learning (https://en.wikipedia.org/wiki/Instance-based_learning).

The complete algorithm

- Choose or invent a kernel function $k(\mathbf{x}, \mathbf{x}') = \phi(\mathbf{x})^T \phi(\mathbf{x}')$ for the $n \times n$ kernel (pairwise-similarity) matrix K
- Compute K for solving $K\mathbf{u} = \lambda\mathbf{u}$
- Center the kernel matrix by

$$K_{centered} = K - \mathbf{1}_n K - K \mathbf{1}_n + \mathbf{1}_n K \mathbf{1}_n$$

Where $\mathbf{1}_n$ is an $n \times n$ matrix with all $\frac{1}{n}$

- Collect top- k eigenvectors \mathbf{u}

The eigenvectors are samples already projected onto principal axes of feature space

Why center the kernel matrix ?

It is similar to the standardization process in *standard* PCA.

But when doing kernel PCA, we do not know whether the ϕ -transformed data is zero-mean, hence the step becomes necessary.

How to center the kernel matrix ?

For each element K_{c-ij} of the centered kernel matrix K_c

TODO (binomial expansion to the projected means)

ref-1 (http://www.ics.uci.edu/~welling/classnotes/papers_class/Kernel-PCA.pdf), **ref-2**
(http://www.cs.haifa.ac.il/~rita/uml_course/lectures/KPCA.pdf)

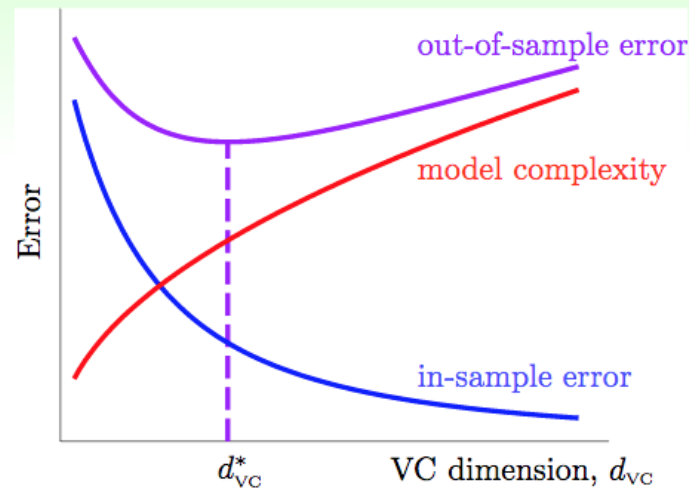
Why \mathbf{u} is projected samples ?

The projected sample is:

$$\begin{aligned}\mathbf{v}^T \phi(X)^T &= \frac{1}{\sqrt{\lambda}} \mathbf{u}^T \phi(X) \phi(X)^T = \frac{1}{\sqrt{\lambda}} \mathbf{u}^T K \\ \implies \phi(X) \mathbf{v} &= \frac{1}{\sqrt{\lambda}} K \mathbf{u} = \sqrt{\lambda} \mathbf{u}\end{aligned}$$

Apply transpose to both side since K is symmetric.

Linear Model First



- tempting sin: use \mathcal{H}_{1126} , low $E_{in}(g_{1126})$ to fool your boss
—**really? :-(a dangerous path of no return**
- safe route: \mathcal{H}_1 first
 - if $E_{in}(g_1)$ good enough, **live happily thereafter :-)**
 - otherwise, move right of the curve
with nothing lost except 'wasted' computation

linear model first:
simple, efficient, **safe**, and **workable!**

The secret of Radial Basis Function (RBF) kernel

- <https://stats.stackexchange.com/questions/80398> (<https://stats.stackexchange.com/questions/80398>)
- <https://stats.stackexchange.com/questions/131138>
(<https://stats.stackexchange.com/questions/131138>)
- <https://stats.stackexchange.com/questions/172554>
(<https://stats.stackexchange.com/questions/172554>)
- <https://math.stackexchange.com/questions/276707>
(<https://math.stackexchange.com/questions/276707>)

Reference

- <https://www.cs.utah.edu/~piyush/teaching/15-9-slides.pdf>
(<https://www.cs.utah.edu/~piyush/teaching/15-9-slides.pdf>)
- 李政軒 PCA & kPCA (<https://www.youtube.com/watch?v=G2NRnh7W4NQ>)
- 林軒田 ML Foundation, Nonlinear Transform (<https://www.youtube.com/watch?v=8pQ06pku1xA>)