

Making Linear Models Nonlinear via Kernel Methods

Piyush Rai

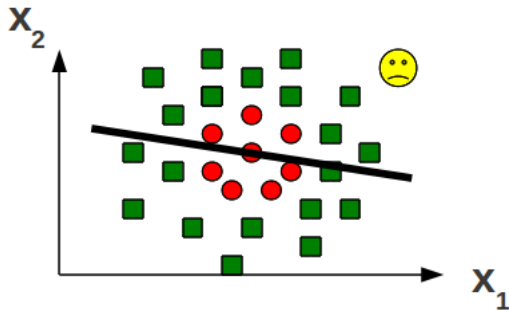
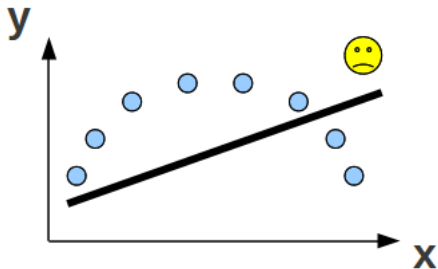
Introduction to Machine Learning (CS771A)

September 6, 2018



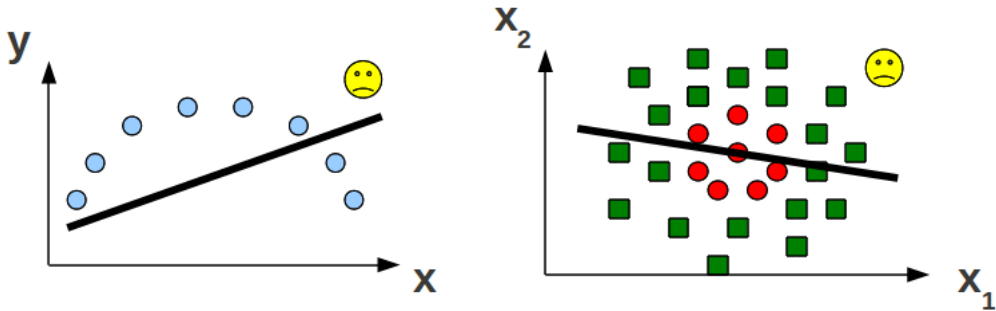
Linear Models

- Nice and interpretable but can't learn “difficult” nonlinear patterns



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- So, are linear models useless for such problems?

Linear Models for Nonlinear Problems!

- Consider the following one-dimensional inputs from two classes



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- Can't separate using a linear hyperplane



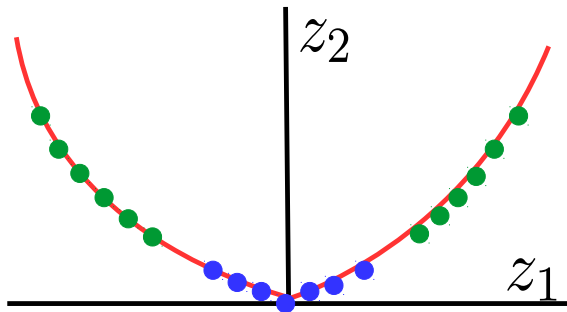
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- Consider mapping each x to two-dimensions as $x \rightarrow \mathbf{z} = [z_1, z_2] = [x, x^2]$



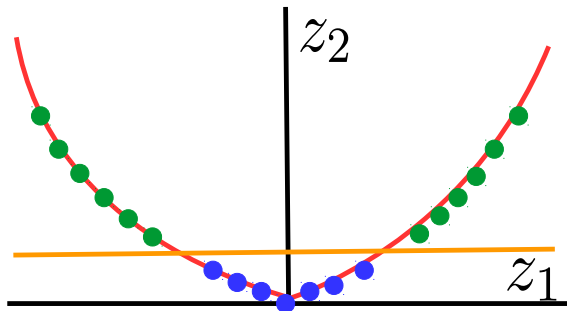
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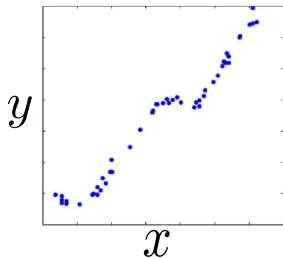
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- Data now becomes linearly separable in the two-dimensional space

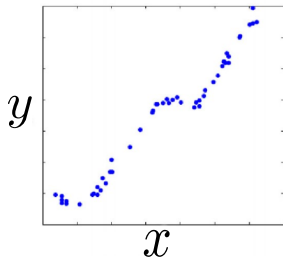
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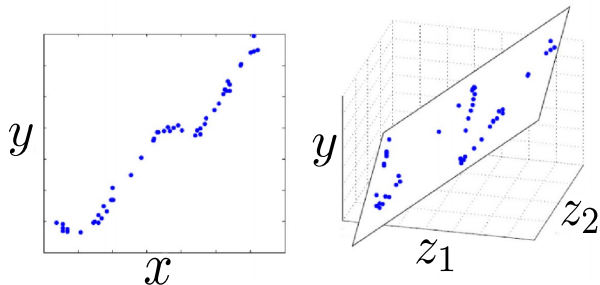


- Linear regression won't work well



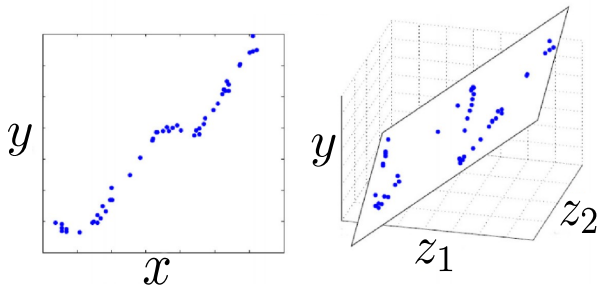
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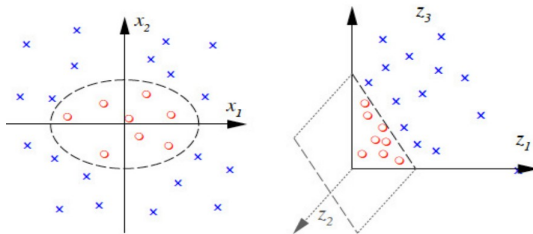
- Now we can fit a linear regression model in two-dimensional input space

Linear Models for Nonlinear Problems!

- Essentially, can use some function ϕ to map/transform inputs to a “nice” space

$$\phi : \mathbb{R}^2 \rightarrow \mathbb{R}^3$$

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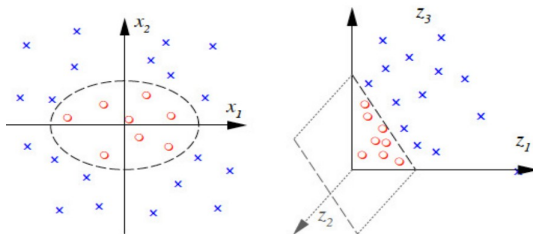


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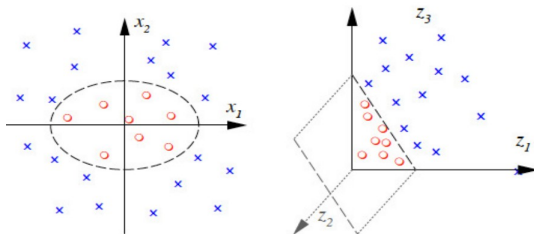


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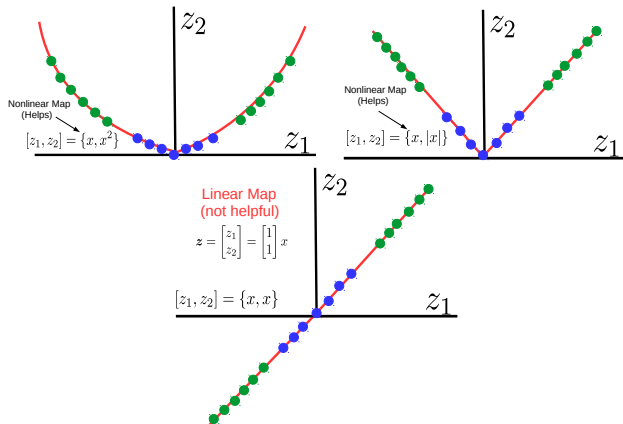
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- .. and then happily apply a linear model in the new space!
- Linear in the new space but nonlinear in the original space!

Not Every Mapping is Helpful

- Not every mapping helps in learning nonlinear patterns. Must at least be nonlinear!
- For the nonlinear classification problem we saw earlier, consider some possible mappings



How to get these “good” (nonlinear) mappings?

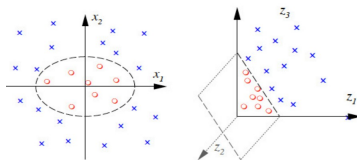
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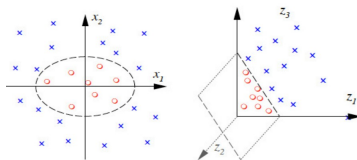
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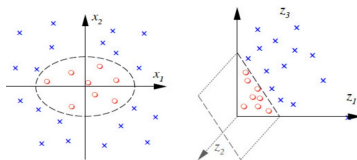
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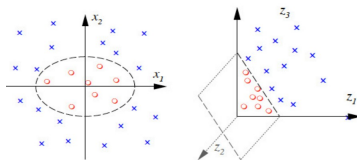


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- Thankfully, not always. For example, when using kernels, you get these for **(almost) free**
 - A kernel defines an “**implicit**” mapping for the data

Kernels as (Implicit) Feature Maps

- Consider two data points $\mathbf{x} = \{x_1, x_2\}$ and $\mathbf{z} = \{z_1, z_2\}$ (each in 2 dims)



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- We didn't need to pre-define/compute the mapping ϕ to compute $k(\mathbf{x}, \mathbf{z})$
- We can simply use the definition of the kernel - $(\mathbf{x}^\top \mathbf{z})^2$ in this case
- Also, evaluating $k(\mathbf{x}, \mathbf{z}) = \phi(\mathbf{x})^\top \phi(\mathbf{z})$ is almost **as fast as computing the inner product $\mathbf{x}^\top \mathbf{z}$**

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$$\phi : \mathcal{X} \rightarrow \mathcal{F}$$

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 - No. The function k must satisfy **Mercer's Condition**



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 - The function k is p.s.d. if the following holds

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.. for all functions f that are “square integrable”, i.e., $\int f(\mathbf{x})^2 d\mathbf{x} < \infty$



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 - $k(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z}) + k_2(\mathbf{x}, \mathbf{z})$: 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



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 - $k(\mathbf{x}, \mathbf{z}) = k_1(\mathbf{x}, \mathbf{z}) k_2(\mathbf{x}, \mathbf{z})$: direct product
 - Kernels can also be constructed by composing these rules



Some Examples of Kernel Functions

- Linear (trivial) Kernel:

$$k(\mathbf{x}, \mathbf{z}) = \mathbf{x}^\top \mathbf{z} \text{ (mapping function } \phi \text{ is identity)}$$



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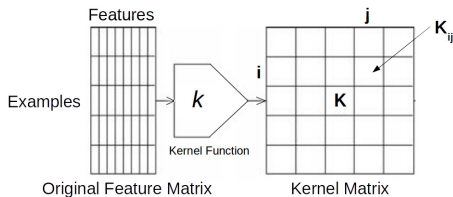


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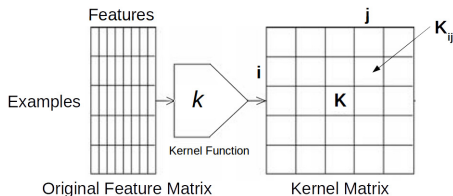


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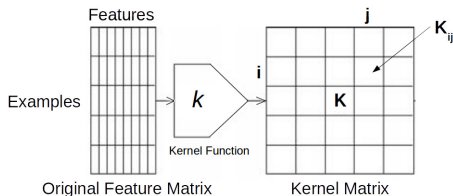


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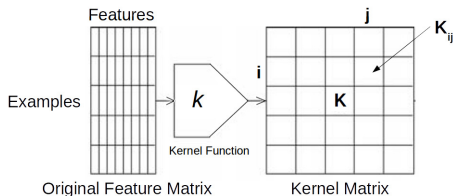
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 - Let's look at two examples: Kernelized SVM and Kernelized Ridge Regression



Example 1:

Kernel (Nonlinear) SVM



Kernelized SVM Training

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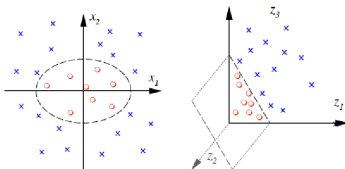
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- Prediction for a new test example \mathbf{x} (assume $b = 0$)

$$y = \text{sign}(\mathbf{w}^\top \phi(\mathbf{x})) = \text{sign}\left(\sum_{n=1}^N \alpha_n y_n \phi(\mathbf{x}_n)^\top \phi(\mathbf{x})\right) = \text{sign}\left(\sum_{n=1}^N \alpha_n y_n k(\mathbf{x}_n, \mathbf{x})\right)$$

- Note: \mathbf{w} can be stored explicitly as a vector only if the feature map $\phi(\cdot)$ can be explicitly written
- In general, kernelized SVMs have to store the training data (at least the support vectors for which α_n 's are nonzero) even at the test time
- Thus the prediction time cost of kernel SVM scales linearly in N
- For unkernelized version $\mathbf{w} = \sum_{n=1}^N \alpha_n y_n \mathbf{x}_n$ can be computed and stored as a $D \times 1$ vector. Thus training data need not be stored and the prediction cost is constant w.r.t. N ($\mathbf{w}^\top \mathbf{x}$ can be computed in $O(D)$ time).

Example 2:

Kernel (Nonlinear) Ridge Regression



Ridge Regression: Revisited

- Recall the ridge regression problem

$$\mathbf{w} = \arg \min_{\mathbf{w}} \sum_{n=1}^N (y_n - \mathbf{w}^\top \mathbf{x}_n)^2 + \lambda \mathbf{w}^\top \mathbf{w}$$



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- Note: $\mathbf{w} = \sum_{n=1}^N \alpha_n \mathbf{x}_n$ is known as “dual” form of ridge regression solution. However, so far it is still a linear model. But now it is easily kernelizable.

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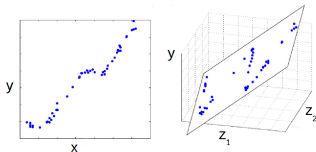
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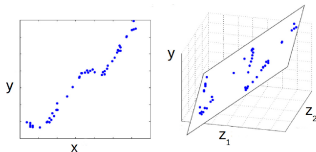
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- Note: Just as in kernel SVM, prediction cost scales in N

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- Kernel methods use a “fixed” set of basis functions or “landmarks”. The basis functions are the training **data points themselves**; also see the next slide.

Kernels: Viewed as Defining Fixed Basis Functions

- Consider each row (or column) of the $N \times N$ kernel matrix (it's symmetric)

$$K = \begin{bmatrix} k(x_1, x_1) & k(x_1, x_2) & \cdots & k(x_1, x_n) \\ k(x_2, x_1) & k(x_2, x_2) & \cdots & k(x_2, x_n) \\ \vdots & \vdots & \ddots & \vdots \\ k(x_n, x_1) & k(x_n, x_2) & \cdots & k(x_n, x_n) \end{bmatrix}$$



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- In general, the set of basis points or landmarks can be **any set of points** (not necessarily the data points) and **can even be learned** (which is what Adaptive Basis Function methods basically do).

Learning with Kernels: Some Aspects (Contd.)

- Storage/computational efficiency can be a bottleneck when using kernels



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- There is a huge literature on **speeding up kernel methods**
 - Approximating the kernel matrix using a set of kernel-derived new features
 - Identifying a small set of landmark points in the training data
 - .. and a lot more



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- All the computations remain as efficient as in the original space
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 - Strings (string kernels): DNA matching, text classification, etc.



Kernels: Concluding Notes

- Kernels give a modular way to learn nonlinear patterns using linear models
 - All you need to do is replace the inner products with the kernel
- All the computations remain as efficient as in the original space
- A very general notion of similarity: Can define similarities between objects even though they can't be represented as vectors. Many kernels are tailor-made for specific types of data
 - Strings (string kernels): DNA matching, text classification, etc.
 - Trees (tree kernels): Comparing parse trees of phrases/sentences

