

Kernel Methods

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Topics in Kernel Methods

1. Kernel Methods vs Linear Models/Neural Networks
2. Stored Sample Methods
3. Kernel Functions
4. Dual Representations
5. Constructing Kernels
6. Extension to Symbolic Inputs
7. Fisher Kernel

Kernel Methods vs Linear Models/Neural Networks

- Linear parametric models for regression and classification have the form $y(x, w)$
 - During learning phase we either get a maximum likelihood estimate of w or a posterior distribution of w
 - Training data is then discarded
 - Prediction based only on vector w
- This is true of Neural networks as well
- Another class of methods use the training samples or a subset of them

Memory-Based Methods

- Training data points are used in prediction phase
- Examples of such methods
 - Parzen probability density model
 - Linear combination of kernel functions centered on each training data point
 - Nearest neighbor classification
- These are memory-based methods
- Require a metric to be defined
- Fast to train, slow to predict

Kernel Functions

- Many linear parametric models can be re-cast into equivalent dual representations where predictions are based on a kernel function evaluated at training points
- Kernel function is given by

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

- where $\phi(\mathbf{x})$ is a fixed nonlinear feature space mapping (basis function)
- Kernel is a symmetric function of its arguments
$$k(\mathbf{x}, \mathbf{x}') = k(\mathbf{x}', \mathbf{x})$$
- Kernel function can be interpreted as the similarity of \mathbf{x} and \mathbf{x}'
- Simplest is identity mapping in feature space $\phi(\mathbf{x}) = \mathbf{x}$
 - In which case $k(\mathbf{x}, \mathbf{x}') = \mathbf{x}^T \mathbf{x}'$
 - Called Linear Kernel

Kernel Trick (or Kernel Substitution)

- Formulated as inner product allows extending well-known algorithms
 - by using the kernel trick
- Basic idea of kernel trick
 - If an input vector \mathbf{x} appears only in the form of scalar products then we can replace scalar products with some other choice of kernel
- Used widely
 - in support vector machines
 - in developing non-linear variant of PCA
 - In kernel Fisher discriminant

Other Forms of Kernel Functions

- Function of difference between arguments

$$k(\mathbf{x}, \mathbf{x}') = k(\mathbf{x} - \mathbf{x}')$$

- Called *stationary* kernel since invariant to translation in space

- *Homogeneous* kernels, also known as *radial basis functions*

$$k(\mathbf{x}, \mathbf{x}') = k(\|\mathbf{x} - \mathbf{x}'\|)$$

- Depend only on the magnitude of the distance between arguments
- Note that the kernel function is a scalar value while \mathbf{x} is an M -dimensional vector

For these to be valid kernel functions they should be shown to have the property

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

Dual Representation

- Linear models for regression and classification can be reformulated in terms of a dual representation
 - In which kernel function arises naturally
- Plays important role in SVMs
- Consider linear regression model
 - whose parameters are determined by minimizing *regularized sum-of-squares* error function

$$J(w) = \frac{1}{2} \sum_{n=1}^N \{w^T \phi(x_n) - t_n\}^2 + \frac{\lambda}{2} w^T w$$

where $w = (w_0, \dots, w_{M-1})^T$, $\phi = (\phi_0, \dots, \phi_{M-1})^T$

we have N samples $\{x_1, \dots, x_N\}$

λ is the regularization coefficient

ϕ is the set of M
basis functions
or feature vector

- Minimum obtained by setting gradient of $J(w)$ wrt w equal to zero

Solution for \mathbf{w} as a linear combination of $\phi(\mathbf{x}_n)$

- By equating derivative $J(\mathbf{w})$ wrt \mathbf{w} to zero and solving for \mathbf{w} we get

$$\begin{aligned}\mathbf{w} &= -\frac{1}{\lambda} \sum_{n=1}^N \{ \mathbf{w}^T \phi(\mathbf{x}_n) - t_n \} \phi(\mathbf{x}_n) \\ &= \sum_{n=1}^N a_n \phi(\mathbf{x}_n) \\ &= \Phi^T \mathbf{a}\end{aligned}$$

- Solution for \mathbf{w} is a linear combination of vectors $\phi(\mathbf{x}_n)$ whose coefficients are functions of \mathbf{w} where

- Φ is the design matrix whose n^{th} row is given by $\phi(\mathbf{x}_n)^T$

$$\Phi = \begin{bmatrix} \phi_0(x_1) & \cdot & \cdot & \phi_{M-1}(x_1) \\ \vdots & & & \vdots \\ \phi_0(x_n) & \cdot & \cdot & \phi_{M-1}(x_n) \\ \vdots & & & \vdots \\ \phi_0(x_N) & \cdot & \cdot & \phi_{M-1}(x_N) \end{bmatrix} \text{ is a } N \times M \text{ matrix}$$

- Vector $\mathbf{a} = (a_1, \dots, a_N)^T$ with the definition

$$a_n = -\frac{1}{\lambda} \{ \mathbf{w}^T \phi(\mathbf{x}_n) - t_n \}$$

Transformation from \mathbf{w} to \mathbf{a}

- Thus we have $\mathbf{w} = \Phi^T \mathbf{a}$
- Instead of working with parameter vector \mathbf{w} we can reformulate least squares algorithm in terms of parameter vector \mathbf{a}
 - giving rise to dual representation
- We will see that although the definition of \mathbf{a} still includes \mathbf{w}

$$a_n = -\frac{1}{\lambda} \left\{ \mathbf{w}^T \phi(\mathbf{x}_n) - t_n \right\}$$

It can be eliminated by the use of the kernel function

Gram Matrix and Kernel Function

- Define the Gram matrix $K = \Phi \Phi^T$ an $N \times N$ matrix, with elements

Note: $N \times M$ times $M \times N$

$$K_{nm} = \phi(x_n)^T \phi(x_m) = k(x_n, x_m)$$

- where we introduce the kernel function $k(x, x') = \phi(x)^T \phi(x')$

$$K = \begin{bmatrix} k(x_1, x_1) & \cdot & \cdot & k(x_1, x_N) \\ \cdot & & & \\ \cdot & & & \\ k(x_N, x_1) & & & k(x_N, x_N) \end{bmatrix}$$

Gram Matrix Definition:
Given N vectors, it is the matrix of all inner products

- Notes:
 - Φ is $N \times M$ and K is $N \times N$
 - K is a matrix of similarities of pairs of samples (thus it is symmetric)

Error Function in Terms of Gram Matrix of Kernel

- Sum of squares Error Function is

$$J(\mathbf{w}) = \frac{1}{2} \sum_{n=1}^N \left\{ \mathbf{w}^T \phi(\mathbf{x}_n) - t_n \right\}^2 + \frac{\lambda}{2} \mathbf{w}^T \mathbf{w}$$

- Substituting $\mathbf{w} = \Phi^T \mathbf{a}$ into $J(\mathbf{w})$ gives

$$J(\mathbf{w}) = \frac{1}{2} \mathbf{a}^T \Phi \Phi^T \Phi \Phi^T \mathbf{a} - \mathbf{a}^T \Phi \Phi^T \mathbf{t} + \frac{1}{2} \mathbf{t}^T \mathbf{t} + \frac{\lambda}{2} \mathbf{a}^T \Phi \Phi^T \mathbf{a}$$

where $\mathbf{t} = (t_1, \dots, t_N)^T$

- Sum of squares error function is written in terms of Gram matrix as

$$J(\mathbf{a}) = \frac{1}{2} \mathbf{a}^T \mathbf{K} \mathbf{K} \mathbf{a} - \mathbf{a}^T \mathbf{K} \mathbf{t} + \frac{1}{2} \mathbf{t}^T \mathbf{t} + \frac{\lambda}{2} \mathbf{a}^T \mathbf{K} \mathbf{a}$$

- Solving for \mathbf{a} by combining $\mathbf{w} = \Phi^T \mathbf{a}$ and $a_n = -\frac{1}{\lambda} \left\{ \mathbf{w}^T \phi(\mathbf{x}_n) - t_n \right\}$

$$\mathbf{a} = (\mathbf{K} + \lambda \mathbf{I}_N)^{-1} \mathbf{t}$$

Solution for \mathbf{a} can be expressed as a linear combination of elements of $\phi(\mathbf{x})$ whose coefficients are entirely in terms of kernel $k(\mathbf{x}, \mathbf{x}')$ from which we can recover original formulation in terms of parameters \mathbf{w}

Prediction Function

- Prediction for new input \mathbf{x}

- We can write $\mathbf{a} = (\mathbf{K} + \lambda \mathbf{I}_N)^{-1} \mathbf{t}$ by combining $\mathbf{w} = \Phi^T \mathbf{a}$ and

$$a_n = -\frac{1}{\lambda} \left\{ \mathbf{w}^T \phi(\mathbf{x}_n) - t_n \right\}$$

- Substituting back into linear regression model,

$$y(\mathbf{x}) = \mathbf{w}^T \phi(\mathbf{x})$$

$$= \mathbf{a}^T \Phi \phi(\mathbf{x})$$

$$= \mathbf{k}(\mathbf{x})^T (\mathbf{K} + \lambda \mathbf{I}_N)^{-1} \mathbf{t} \text{ where } \mathbf{k}(\mathbf{x}) \text{ has elements } k_n(\mathbf{x}) = k(\mathbf{x}_n, \mathbf{x})$$

- Prediction is a linear combination of the target values from the training set.

Advantage of Dual Representation

- Solution for \mathbf{a} is expressed entirely in terms of kernel function $k(\mathbf{x}, \mathbf{x}')$
- Once we get \mathbf{a} we can recover \mathbf{w} as linear combination of elements of $\phi(\mathbf{x})$ using $\mathbf{w} = \Phi^t \mathbf{a}$
- In parametric formulation, solution is $\mathbf{w}_{ML} = (\Phi^T \Phi)^{-1} \Phi^T \mathbf{t}$
 - Instead of inverting an $M \times M$ matrix we are inverting an $N \times N$ matrix— an apparent disadvantage
- But, advantage of dual formulation is that we can work with kernel function $k(\mathbf{x}, \mathbf{x}')$ and therefore
 - avoid working with a feature vector $\phi(\mathbf{x})$ and
 - problems associated with very high or infinite dimensionality of \mathbf{x}

Constructing Kernels

- To exploit kernel substitution need valid kernel functions
- First Method
 - choose a feature space mapping $\phi(x)$ and use it to find corresponding kernel
 - One-dimensional input space

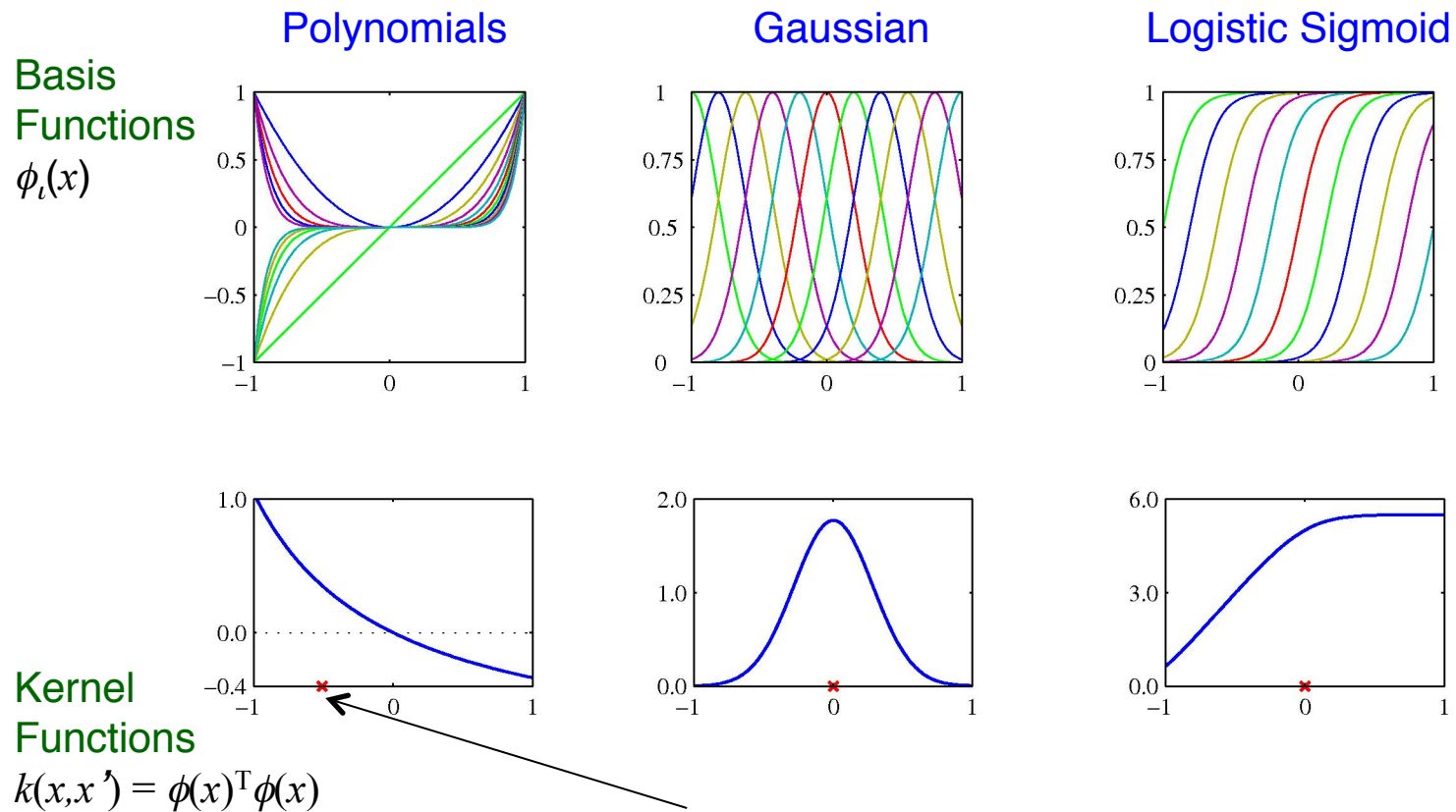
$$k(x, x') = \phi(x)^T \phi(x')$$

$$= \sum_{i=1}^M \phi_i(x) \phi_i(x')$$

- where $\phi(x)$ are basis functions such as polynomial
- For each i we choose $\phi_i = x^i$

Construction of Kernel Functions from basis functions

One-dimensional input space



Red cross is x'

Second Method: Direct Construction of Kernels

- Function we choose has to correspond to a scalar product in some (perhaps infinite dimensional) space
- Consider kernel function $k(\mathbf{x}, \mathbf{z}) = (\mathbf{x}^T \mathbf{z})^2$
 - In two dimensional space

$$\begin{aligned} k(\mathbf{x}, \mathbf{z}) &= (\mathbf{x}^T \mathbf{z})^2 = (x_1 z_1 + x_2 z_2)^2 \\ &= x_1^2 z_1^2 + 2x_1 z_1 x_2 z_2 + x_2^2 z_2^2 \\ &= (x_1^2, \sqrt{2}x_1 x_2, x_2^2)(z_1^2, \sqrt{2}z_1 z_2, z_2^2)^T \\ &= \phi(\mathbf{x})^T \phi(\mathbf{z}) \end{aligned}$$
 - Feature mapping takes the form $\phi(x) = (x_1^2, \sqrt{2}x_1 x_2, x_2^2)$
 - Comprises of all second order terms with a specific weighting
 - Inner product needs computing six feature values and $3 \times 3 = 9$ multiplications
 - Kernel function $k(x, z)$ has 2 multiplications and a squaring
- By considering $(\mathbf{x}^T \mathbf{z} + c)^2$ we get constant, linear, second order terms
- By considering $(\mathbf{x}^T \mathbf{z} + c)^M$ we get all powers of \mathbf{x} (monomials)

Testing whether a function is a valid kernel

- Without having to construct the function $\phi(\mathbf{x})$ explicitly
- Necessary and sufficient condition for a function $k(\mathbf{x}, \mathbf{x}')$ to be a kernel is
 - Gram matrix K , whose elements are given by $k(\mathbf{x}_n, \mathbf{x}_m)$ is positive semi-definite for all possible choices of the set $\{\mathbf{x}_n\}$
 - Positive semi-definite is not the same thing as a matrix whose elements are non-negative
 - It means $\mathbf{z}^T K \mathbf{z} \geq 0$ for non-zero vectors \mathbf{z} with real entries
 i.e., $\sum_n \sum_m K_{nm} z_n z_m \geq 0$ for any real numbers z_n, z_m
 - Mercer's theorem: any continuous, symmetric, positive semi-definite kernel function $k(x, y)$ can be expressed as a dot product in a high-dimensional space
- New kernels can be constructed from simpler kernels as building blocks

Techniques for Constructing Kernels

- Given valid kernels $k_1(x, x')$ and $k_2(x, x')$ the following new kernels will be valid

- $k(x, x') = ck_1(x, x')$
- $k(x, x') = f(x)k_1(x, x')f(x')$
- $k(x, x') = q(k_1(x, x'))$
- $k(x, x') = \exp(k_1(x, x'))$
- $k(x, x') = k_1(x, x') + k_2(x, x')$
- $k(x, x') = k_1(x, x')k_2(x, x')$
- $k(x, x') = k_3(\phi(x) \cdot \phi(x'))$
- $k(x, x') = x^T A x'$
- $k(x, x') = k_a(x_a, x'_a) + k_b(x_b, x'_b)$
- $k(x, x') = k_a(x_a, x'_a)k_b(x_b, x'_b)$

Where

$f(\cdot)$ is any function

$q(\cdot)$ is a polynomial with non-negative coefficients

$\phi(x)$ is a function from x to \mathbb{R}^M

k_3 is a valid kernel in \mathbb{R}^M

A is a symmetric positive semidefinite matrix

x_a and x_b are variables with $x = (x_a, x_b)$

k_a and k_b are valid kernel functions

Kernels appropriate for specific applications

- Requirements for $k(\mathbf{x}, \mathbf{x}')$
 - It is symmetric
 - Its Gram matrix is positive semidefinite
 - It expresses the appropriate similarity between \mathbf{x} and \mathbf{x}' for the intended application

Gaussian Kernel

- Commonly used kernel is

$$k(\mathbf{x}, \mathbf{x}') = \exp(-\|\mathbf{x} - \mathbf{x}'\|^2 / 2\sigma^2)$$

- It is seen as a valid kernel by expanding the square

$$\|\mathbf{x} - \mathbf{x}'\|^2 = \mathbf{x}^T \mathbf{x} + (\mathbf{x}')^T \mathbf{x}' - 2\mathbf{x}^T \mathbf{x}'$$

- To give

$$k(\mathbf{x}, \mathbf{x}') = \exp(-\mathbf{x}^T \mathbf{x} / 2\sigma^2) \exp(-(\mathbf{x}')^T \mathbf{x}' / 2\sigma^2) \exp(\mathbf{x}^T \mathbf{x}' / \sigma^2)$$

- From kernel construction rules 2 and 4

- together with validity of linear kernel $k(\mathbf{x}, \mathbf{x}') = \mathbf{x}^T \mathbf{x}'$

- Can be extended to non-Euclidean distances

$$k(\mathbf{x}, \mathbf{x}') = \exp \{(-1/2\sigma^2)[\kappa(\mathbf{x}, \mathbf{x}') + \kappa(\mathbf{x}', \mathbf{x}') - 2\kappa(\mathbf{x}, \mathbf{x}')]\}$$

Extension of Kernels to Symbolic Inputs

- Important contribution of kernel viewpoint:
 - Inputs that are symbolic rather than vectors of real numbers
- Kernel functions defined for graphs, sets, strings, text documents
- If A_1 and A_2 are two subsets of objects
 - A simple kernel is
$$k(A_1, A_2) = 2^{|A_1 \cap A_2|}$$
 - where $| \cdot |$ indicates cardinality of set intersection
 - A valid kernel since it can be shown to correspond to an inner product in a feature space

$$A = \{1, 2, 3, 4, 5\}$$

$$A_1 = \{2, 3, 4, 5\}$$

$$A_2 = \{1, 2, 4, 5\}$$

$$A_1 \cap A_2 = \{2, 4, 5\}$$

$$\text{Hence } k(A_1, A_2) = 8$$

What are feature vectors $\phi(A_1)$ and $\phi(A_2)$ such that $\phi(A_1)\phi(A_2)^T = 8$?

Combining Discriminative and Generative Models

- Generative models deal naturally with missing data and with HMM of varying length
- Discriminative models such as SVM have better performance
- Can use a generative model to define a kernel and use kernel in discriminative approach

Kernels based on Generative Models

- Given a generative model $p(\mathbf{x})$ we define a kernel by

$$k(\mathbf{x}, \mathbf{x}') = p(\mathbf{x}) p(\mathbf{x}')$$

- A valid kernel since it is an inner product in the one-dimensional feature space defined by the mapping $p(\mathbf{x})$
- Two inputs \mathbf{x} and \mathbf{x}' are similar if they have high probabilities

Kernel Functions based on Mixture Densities

- Extension to sums of products of different probability distributions

$$k(x, x') = \sum_i p(x | i) p(x' | i) p(i)$$

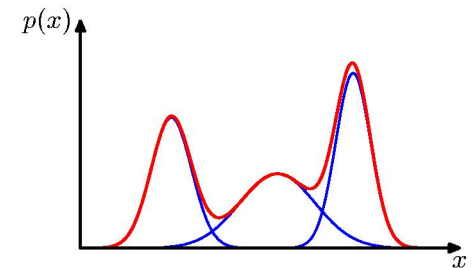
- where $p(i)$ are positive weighting coefficients
- It is a valid kernel based on two rules of kernel construction:

$$k(x, x') = c k_1(x, x') \text{ and } k(x, x') = k_1(x, x') + k_2(x, x')$$

- Two inputs x and x' will give a large value of k , and hence appear similar, if they have a significant probability under a range of different components
- Taking the limit to infinite sum

$$k(x, x') = \int p(x | z) p(x' | z) p(z) dz$$

- where z is a continuous latent variable



Kernels for Sequences

- Data consists of ordered sequences of length L

$$\mathbf{X} = \{x_1, \dots, x_L\}$$

- Generative model for sequences is HMM
 - Hidden states $\mathbf{Z} = \{z_1, \dots, z_L\}$
- Kernel Function for measuring similarity of sequences \mathbf{X} and \mathbf{X}' is

$$k(\mathbf{X}, \mathbf{X}') = \sum_{\mathbf{Z}} p(\mathbf{X} | \mathbf{Z}) p(\mathbf{X}' | \mathbf{Z}) p(\mathbf{Z})$$

- Both observed sequences are generated by same hidden sequence \mathbf{Z}

Fisher Kernel

- Alternative technique for using generative models
 - Used in document retrieval, protein sequences, document recognition
- Consider parametric generative model $p(\mathbf{x}|\theta)$ where θ denotes vector of parameters
- Goal: find kernel that measures similarity of two vectors \mathbf{x} and \mathbf{x}' induced by the generative model
- Define Fisher score as gradient wrt θ

$$\mathbf{g}(\theta, \mathbf{x}) = \nabla_{\theta} \ln p(\mathbf{x} | \theta)$$

A vector of same dimensionality as θ

- Fisher Kernel is

$$k(\mathbf{x}, \mathbf{x}') = \mathbf{g}(\theta, \mathbf{x})^T \mathbf{F}^{-1} \mathbf{g}(\theta, \mathbf{x}')$$

where \mathbf{F} is the Fisher information matrix

Fisher score is more generally the gradient of the log-likelihood

$$\mathbf{F} = \mathbb{E}_{\mathbf{x}} [\mathbf{g}(\theta, \mathbf{x}) \mathbf{g}(\theta, \mathbf{x})^T]$$

Fisher Information Matrix

- Presence of Fisher information matrix causes kernel to be invariant under non-linear parametrization of the density model $\theta \rightarrow \psi(\theta)$
- In practice, infeasible to evaluate Fisher Information Matrix. Instead use the approximation

$$F \approx \frac{1}{N} \sum_{n=1}^N g(\theta, x_n) g(\theta, x_n)^T$$

- This is the covariance matrix of the Fisher scores
- So the Fisher kernel $k(x, x') = g(\theta, x)^T F^{-1} g(\theta, x')$ corresponds to whitening of the Fisher scores
- More simply omit F and use non-invariant kernel

$$k(x, x') = g(\theta, x)^T g(\theta, x')$$

Sigmoidal Kernel

- Provides a link between SVMs and neural networks

$$k(\mathbf{x}, \mathbf{x}') = \tanh(a\mathbf{x}^T \mathbf{x}' + b)$$

- Its Gram matrix is not positive semidefinite
 - But used in practice because it gives SVMs a superficial resemblance to neural networks
- Bayesian neural network with an appropriate prior reduces to a Gaussian process
 - Provides a deeper link between neural networks and kernel methods