

# A Tutorial on Gaussian Processes (or why I don't use SVMs)

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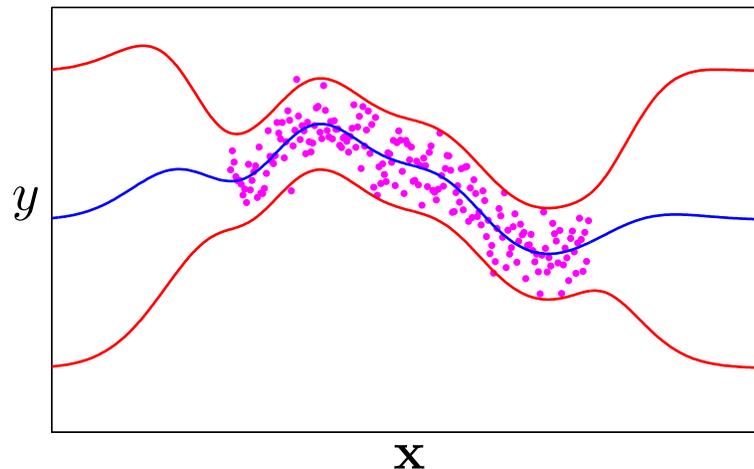
`http://learning.eng.cam.ac.uk/zoubin/`

**MLSS 2011**

# Nonlinear regression

Consider the problem of **nonlinear regression**:

You want to learn a **function**  $f$  with **error bars** from **data**  $\mathcal{D} = \{\mathbf{X}, \mathbf{y}\}$



A **Gaussian process** defines a distribution over functions  $p(f)$  which can be used for Bayesian regression:

$$p(f|\mathcal{D}) = \frac{p(f)p(\mathcal{D}|f)}{p(\mathcal{D})}$$

# Gaussian Processes

A Gaussian process defines a distribution over functions,  $p(f)$ , where  $f$  is a function mapping some input space  $\mathcal{X}$  to  $\mathbb{R}$ .

$$f : \mathcal{X} \rightarrow \mathbb{R}.$$

Notice that  $f$  can be an infinite-dimensional quantity (e.g. if  $\mathcal{X} = \mathbb{R}$ )

Let  $\mathbf{f} = (f(x_1), \dots, f(x_n))$  be an  $n$ -dimensional vector of function values evaluated at  $n$  points  $x_i \in \mathcal{X}$ . Note  $\mathbf{f}$  is a random variable.

**Definition:**  $p(f)$  is a **Gaussian process** if for *any* finite subset  $\{x_1, \dots, x_n\} \subset \mathcal{X}$ , the marginal distribution over that finite subset  $p(\mathbf{f})$  has a multivariate Gaussian distribution.

# Gaussian process covariance functions (kernels)

$p(f)$  is a **Gaussian process** if for *any* finite subset  $\{x_1, \dots, x_n\} \subset \mathcal{X}$ , the marginal distribution over that finite subset  $p(\mathbf{f})$  has a multivariate Gaussian distribution.

Gaussian processes (GPs) are parameterized by a **mean function**,  $\mu(x)$ , and a **covariance function, or kernel**,  $K(x, x')$ .

$$p(f(x), f(x')) = \mathbf{N}(\boldsymbol{\mu}, \boldsymbol{\Sigma})$$

where

$$\boldsymbol{\mu} = \begin{bmatrix} \mu(x) \\ \mu(x') \end{bmatrix} \quad \boldsymbol{\Sigma} = \begin{bmatrix} K(x, x) & K(x, x') \\ K(x', x) & K(x', x') \end{bmatrix}$$

and similarly for  $p(f(x_1), \dots, f(x_n))$  where now  $\boldsymbol{\mu}$  is an  $n \times 1$  vector and  $\boldsymbol{\Sigma}$  is an  $n \times n$  matrix.

# Gaussian process covariance functions

Gaussian processes (GPs) are parameterized by a **mean function**,  $\mu(x)$ , and a **covariance function**,  $K(x, x')$ .

An example covariance function:

$$K(x_i, x_j) = v_0 \exp \left\{ - \left( \frac{|x_i - x_j|}{r} \right)^\alpha \right\} + v_1 + v_2 \delta_{ij}$$

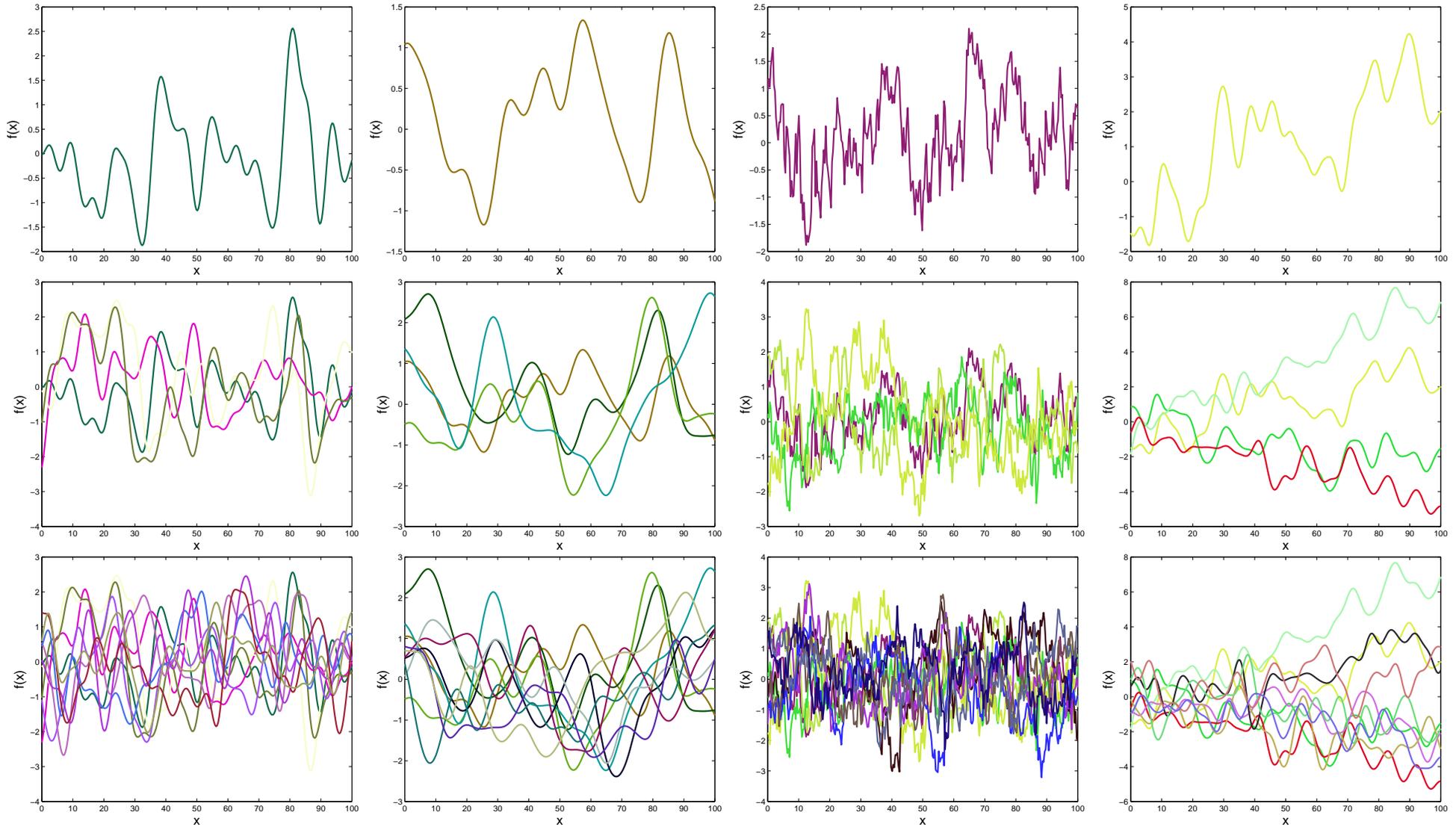
with parameters  $(v_0, v_1, v_2, r, \alpha)$

These kernel parameters are **interpretable** and can be learned from data:

$v_0$	signal variance
$v_1$	variance of bias
$v_2$	noise variance
$r$	lengthscale
$\alpha$	roughness

Once the mean and covariance functions are defined, everything else about GPs follows from the basic rules of probability applied to multivariate Gaussians.

# Samples from GPs with different $K(x, x')$



# Using Gaussian processes for nonlinear regression

Imagine observing a data set  $\mathcal{D} = \{(\mathbf{x}_i, y_i)_{i=1}^n\} = (\mathbf{X}, \mathbf{y})$ .

Model:

$$\begin{aligned}y_i &= f(\mathbf{x}_i) + \epsilon_i \\f &\sim \text{GP}(\cdot|0, K) \\ \epsilon_i &\sim \text{N}(\cdot|0, \sigma^2)\end{aligned}$$

Prior on  $f$  is a GP, likelihood is Gaussian, therefore posterior on  $f$  is also a GP.

We can use this to make **predictions**

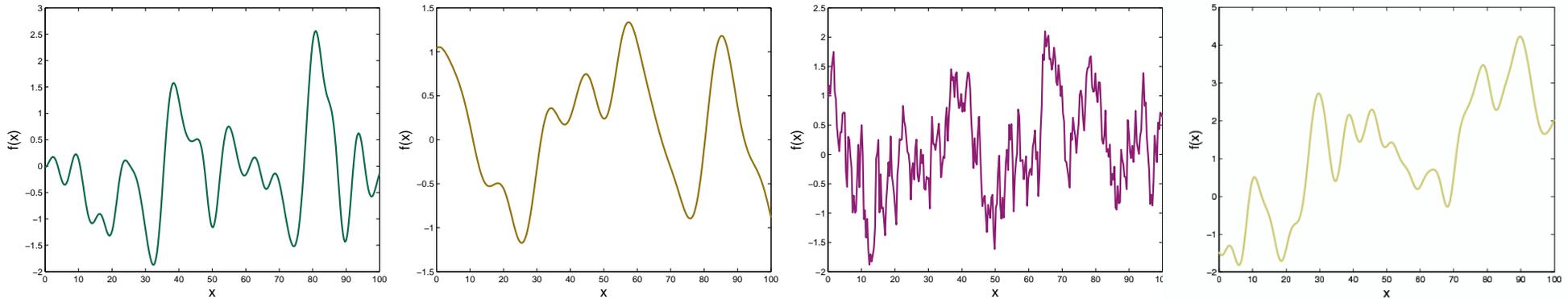
$$p(y_*|\mathbf{x}_*, \mathcal{D}) = \int p(y_*|\mathbf{x}_*, f, \mathcal{D}) p(f|\mathcal{D}) df$$

We can also compute the **marginal likelihood** (evidence) and use this to compare or tune covariance functions

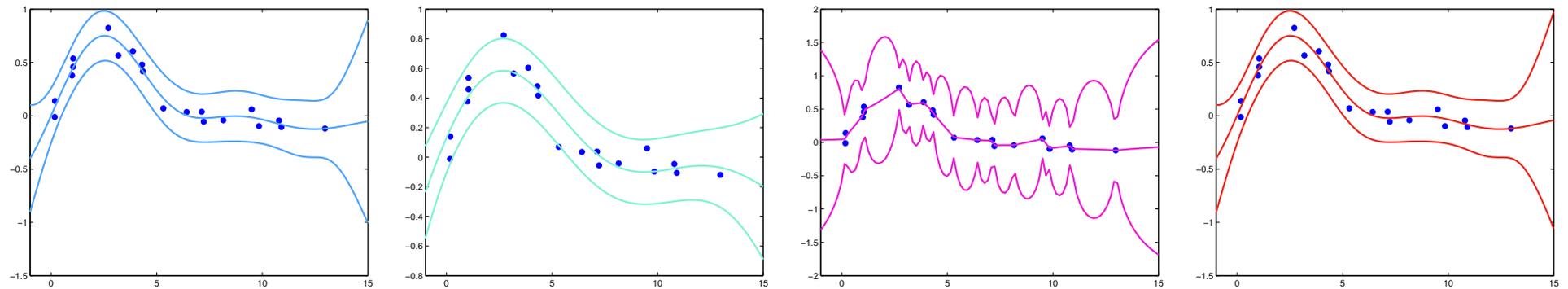
$$p(\mathbf{y}|\mathbf{X}) = \int p(\mathbf{y}|f, \mathbf{X}) p(f) df$$

# Prediction using GPs with different $K(x, x')$

A sample from the prior for each covariance function:



Corresponding predictions, mean with two standard deviations:

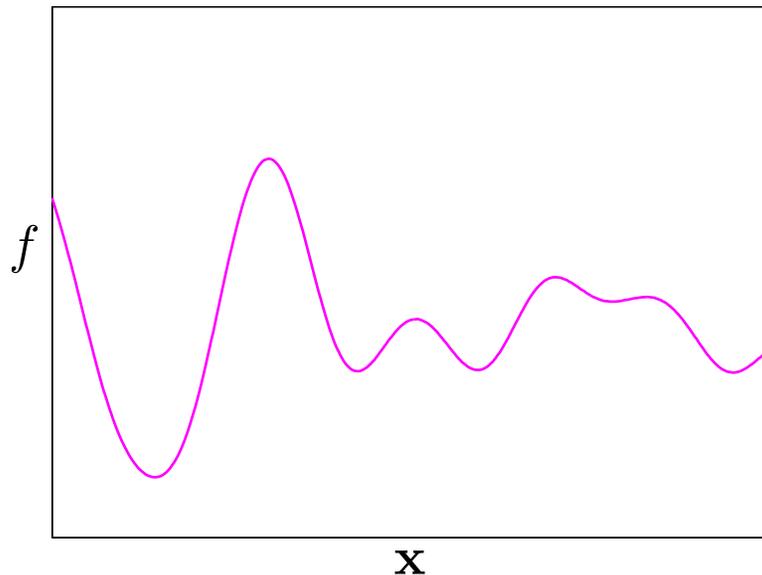


gpdemo

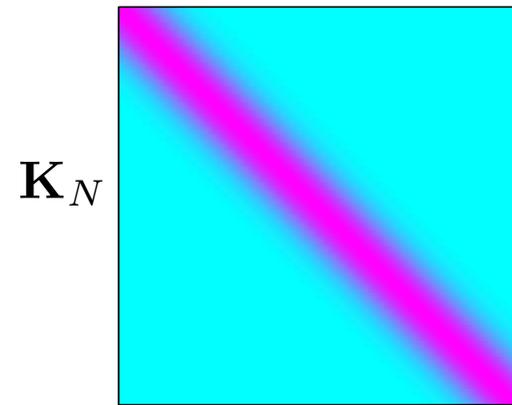
# Gaussian process (GP) priors

**GP**: consistent Gaussian prior on any set of function values  $\mathbf{f} = \{f_n\}_{n=1}^N$ , given corresponding inputs  $\mathbf{X} = \{\mathbf{x}_n\}_{n=1}^N$

one sample function



prior  
 $p(\mathbf{f}|\mathbf{X}) = \mathcal{N}(\mathbf{0}, \mathbf{K}_N)$

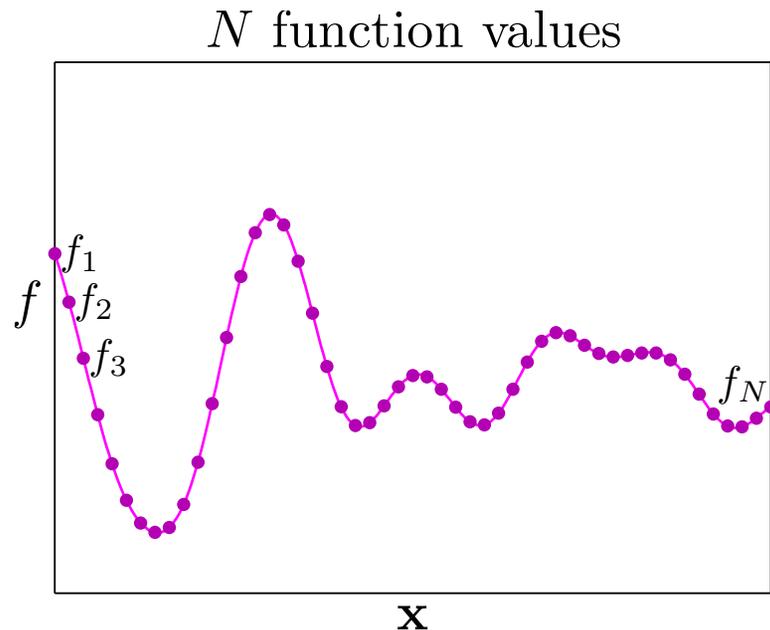


**Covariance**:  $\mathbf{K}_{nn'} = K(\mathbf{x}_n, \mathbf{x}_{n'}; \boldsymbol{\theta})$ , hyperparameters  $\boldsymbol{\theta}$

$$\mathbf{K}_{nn'} = v \exp \left[ -\frac{1}{2} \sum_{d=1}^D \left( \frac{x_n^{(d)} - x_{n'}^{(d)}}{r_d} \right)^2 \right]$$

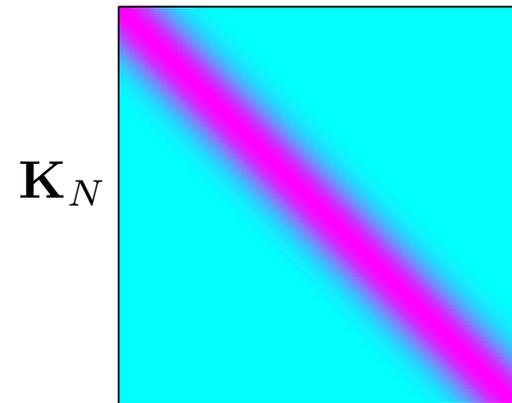
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prior

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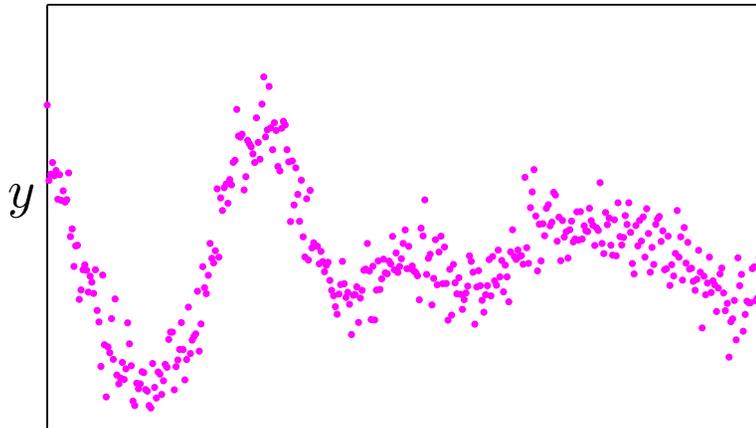
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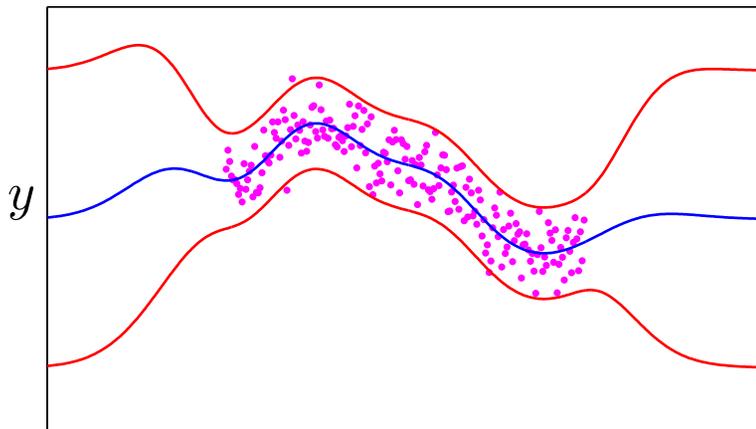
# GP regression

Gaussian observation noise:  $y_n = f_n + \epsilon_n$ , where  $\epsilon_n \sim \mathcal{N}(0, \sigma^2)$

sample data



predictive



marginal likelihood

$$p(\mathbf{y}|\mathbf{X}) = \mathcal{N}(\mathbf{0}, \mathbf{K}_N + \sigma^2\mathbf{I})$$

predictive distribution

$$p(y_*|\mathbf{x}_*, \mathbf{X}, \mathbf{y}) = \mathcal{N}(\mu_*, \sigma_*^2)$$

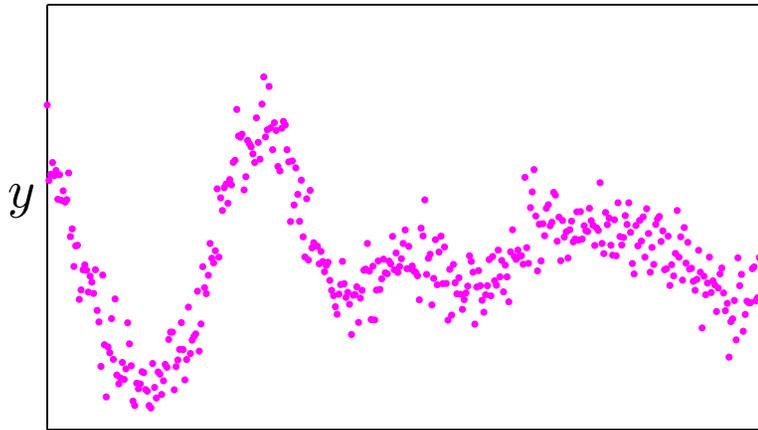
$$\mu_* = \mathbf{K}_{*N}(\mathbf{K}_N + \sigma^2\mathbf{I})^{-1}\mathbf{y}$$

$$\sigma_*^2 = K_{**} - \mathbf{K}_{*N}(\mathbf{K}_N + \sigma^2\mathbf{I})^{-1}\mathbf{K}_{N*} + \sigma^2$$

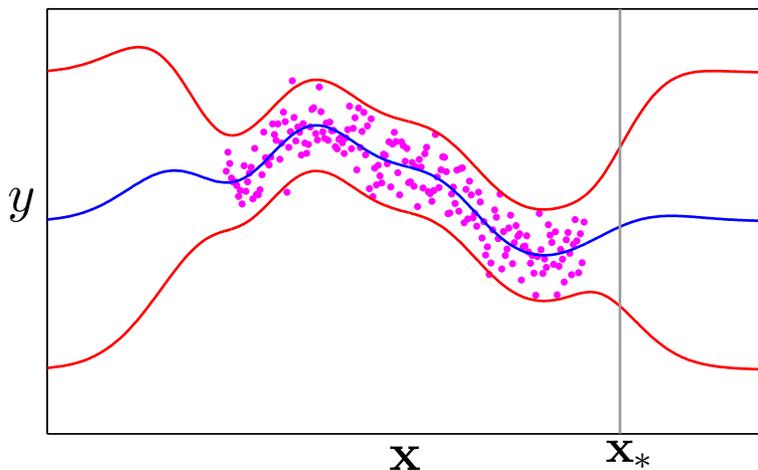
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$$\sigma_*^2 = K_{**} - \mathbf{K}_{*N}(\mathbf{K}_N + \sigma^2\mathbf{I})^{-1}\mathbf{K}_{N*} + \sigma^2$$

# GP learning the kernel

Consider the **covariance function**  $K$  with hyperparameters  $\boldsymbol{\theta} = (v_0, v_1, r_1, \dots, r_d, \alpha)$ :

$$K_{\boldsymbol{\theta}}(\mathbf{x}_i, \mathbf{x}_j) = v_0 \exp \left\{ - \sum_{d=1}^D \left( \frac{|x_i^{(d)} - x_j^{(d)}|}{r_d} \right)^{\alpha} \right\} + v_1$$

Given a data set  $\mathcal{D} = (\mathbf{X}, \mathbf{y})$ , how do we learn  $\boldsymbol{\theta}$ ?

The **marginal likelihood** is a function of  $\boldsymbol{\theta}$

$$p(\mathbf{y}|\mathbf{X}, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{0}, \mathbf{K}_{\boldsymbol{\theta}} + \sigma^2 \mathbf{I})$$

where its log is:

$$\ln p(\mathbf{y}|\mathbf{X}, \boldsymbol{\theta}) = -\frac{1}{2} \ln \det(\mathbf{K}_{\boldsymbol{\theta}} + \sigma^2 \mathbf{I}) - \frac{1}{2} \mathbf{y}^{\top} (\mathbf{K}_{\boldsymbol{\theta}} + \sigma^2 \mathbf{I})^{-1} \mathbf{y} + \text{const}$$

which can be optimized as a function of  $\boldsymbol{\theta}$  and  $\sigma$ .

Alternatively, one can infer  $\boldsymbol{\theta}$  using Bayesian methods, which is more costly but immune to overfitting.

# From linear regression to GPs:

- Linear regression with inputs  $x_i$  and outputs  $y_i$ :  $y_i = \beta_0 + \beta_1 x_i + \epsilon_i$

- Linear regression with  $M$  basis functions:  $y_i = \sum_{m=1}^M \beta_m \phi_m(x_i) + \epsilon_i$

- Bayesian linear regression with basis functions:

$$\beta_m \sim \mathbf{N}(\cdot | 0, \lambda_m) \quad (\text{independent of } \beta_\ell, \forall \ell \neq m), \quad \epsilon_i \sim \mathbf{N}(\cdot | 0, \sigma^2)$$

- Integrating out the coefficients,  $\beta_j$ , we find:

$$E[y_i] = 0, \quad \text{Cov}(y_i, y_j) = K_{ij} \stackrel{\text{def}}{=} \sum_{m=1}^M \lambda_m \phi_m(x_i) \phi_m(x_j) + \delta_{ij} \sigma^2$$

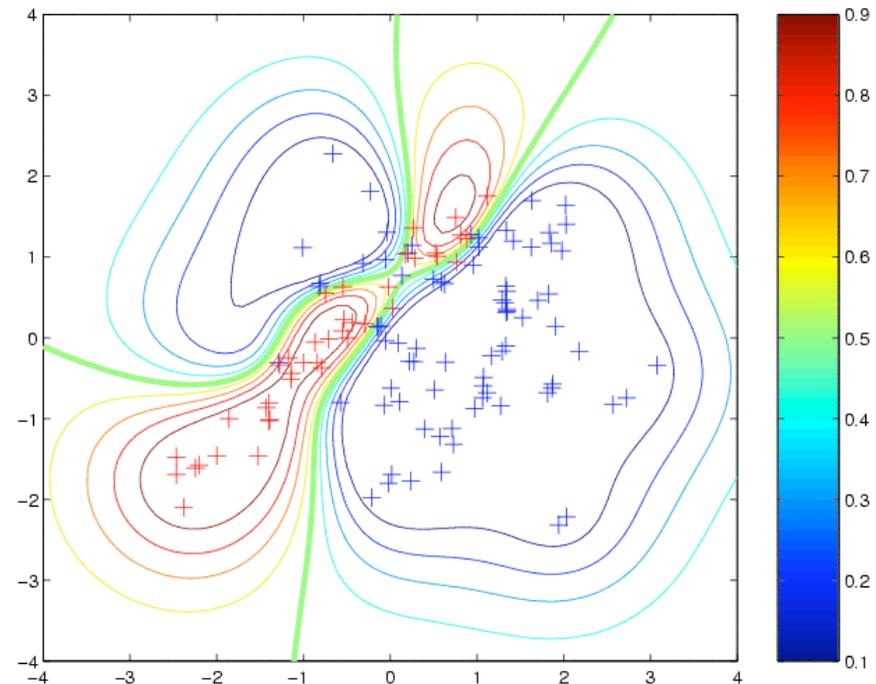
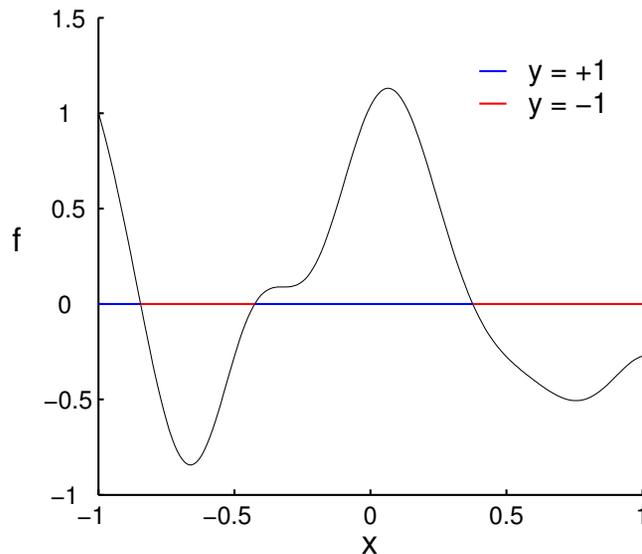
This is a Gaussian process with covariance function  $K(x_i, x_j) = K_{ij}$ .

This GP has a finite number ( $M$ ) of basis functions. Many useful GP kernels correspond to infinitely many basis functions (i.e. infinite-dim feature spaces).

A multilayer perceptron (neural network) with infinitely many hidden units and Gaussian priors on the weights  $\rightarrow$  a GP (Neal, 1996)

# Using Gaussian Processes for Classification

**Binary classification problem:** Given a data set  $\mathcal{D} = \{(\mathbf{x}_i, y_i)\}_{i=1}^n$ , with binary class labels  $y_i \in \{-1, +1\}$ , infer class label probabilities at new points.



There are many ways to relate function values  $f_i = f(\mathbf{x}_i)$  to class probabilities:

$$p(y_i|f_i) = \begin{cases} \frac{1}{1+\exp(-y_i f_i)} & \text{sigmoid (logistic)} \\ \Phi(y_i f_i) & \text{cumulative normal (probit)} \\ \mathbf{H}(y_i f_i) & \text{threshold} \\ \epsilon + (1 - 2\epsilon)\mathbf{H}(y_i f_i) & \text{robust threshold} \end{cases}$$

Non-Gaussian likelihood, so we need to use approximate inference methods (Laplace, EP, MCMC).

# Support Vector Machines

Consider soft-margin Support Vector Machines:

$$\min_{\mathbf{w}} \frac{1}{2} \|\mathbf{w}\|^2 + C \sum_i (1 - y_i f_i)_+$$

where  $(\cdot)_+$  is the hinge loss and  $f_i = f(\mathbf{x}_i) = \mathbf{w} \cdot \mathbf{x}_i + w_0$ . Let's kernelize this:

$$\mathbf{x}_i \rightarrow \phi(\mathbf{x}_i) = k(\cdot, \mathbf{x}_i), \quad \mathbf{w} \rightarrow f(\cdot)$$

By reproducing property:

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

By representer theorem, solution:

$$f(\mathbf{x}) = \sum_i \alpha_i k(\mathbf{x}, \mathbf{x}_i)$$

Defining  $\mathbf{f} = (f_1, \dots, f_N)^T$  note that  $\mathbf{f} = \mathbf{K}\boldsymbol{\alpha}$ , so  $\boldsymbol{\alpha} = \mathbf{K}^{-1}\mathbf{f}$

Therefore the regularizer  $\frac{1}{2} \|\mathbf{w}\|^2 \rightarrow \frac{1}{2} \|f\|_{\mathcal{H}}^2 = \frac{1}{2} \langle f(\cdot), f(\cdot) \rangle_{\mathcal{H}} = \frac{1}{2} \boldsymbol{\alpha}^\top \mathbf{K} \boldsymbol{\alpha} = \frac{1}{2} \mathbf{f}^\top \mathbf{K}^{-1} \mathbf{f}$

So we can rewrite the kernelized SVM loss as:

$$\min_{\mathbf{f}} \frac{1}{2} \mathbf{f}^\top \mathbf{K}^{-1} \mathbf{f} + C \sum_i (1 - y_i f_i)_+$$

# Support Vector Machines and Gaussian Processes

We can write the SVM loss as:

$$\min_{\mathbf{f}} \frac{1}{2} \mathbf{f}^\top \mathbf{K}^{-1} \mathbf{f} + C \sum_i (1 - y_i f_i)_+$$

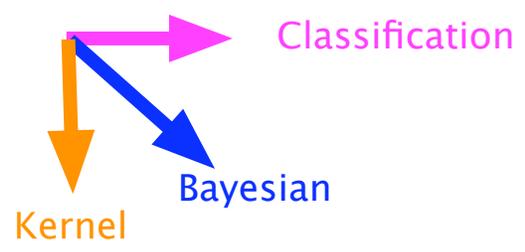
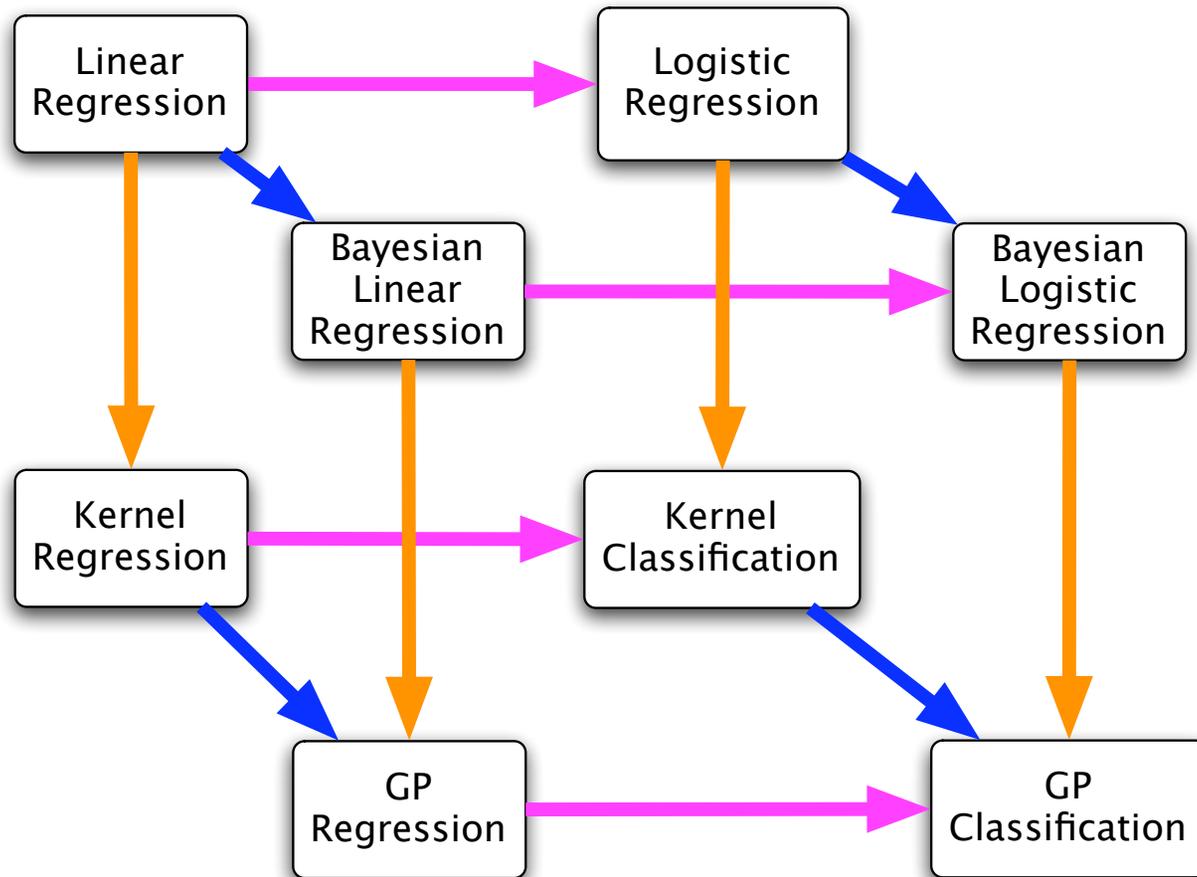
We can write the negative log of a GP likelihood as:  $\frac{1}{2} \mathbf{f}^\top \mathbf{K}^{-1} \mathbf{f} - \sum_i \ln p(y_i | f_i) + c$

Equivalent? No.

With Gaussian processes we:

- Handle **uncertainty** in unknown function  $\mathbf{f}$  by averaging, not minimization.
- Compute  $p(y = +1 | \mathbf{x}) \neq p(y = +1 | \hat{\mathbf{f}}, \mathbf{x})$ .
- Can **learn the kernel parameters** automatically from data, no matter how flexible we wish to make the kernel.
- Can **learn the regularization parameter**  $C$  without cross-validation.
- Can incorporate **interpretable** noise models and priors over functions, and can sample from prior to get intuitions about the model assumptions.
- We can combine **automatic feature selection** with learning using ARD.

# A picture



# Matlab Demo: Gaussian Process Classification

matlab/gpml-matlab/gpml-demo

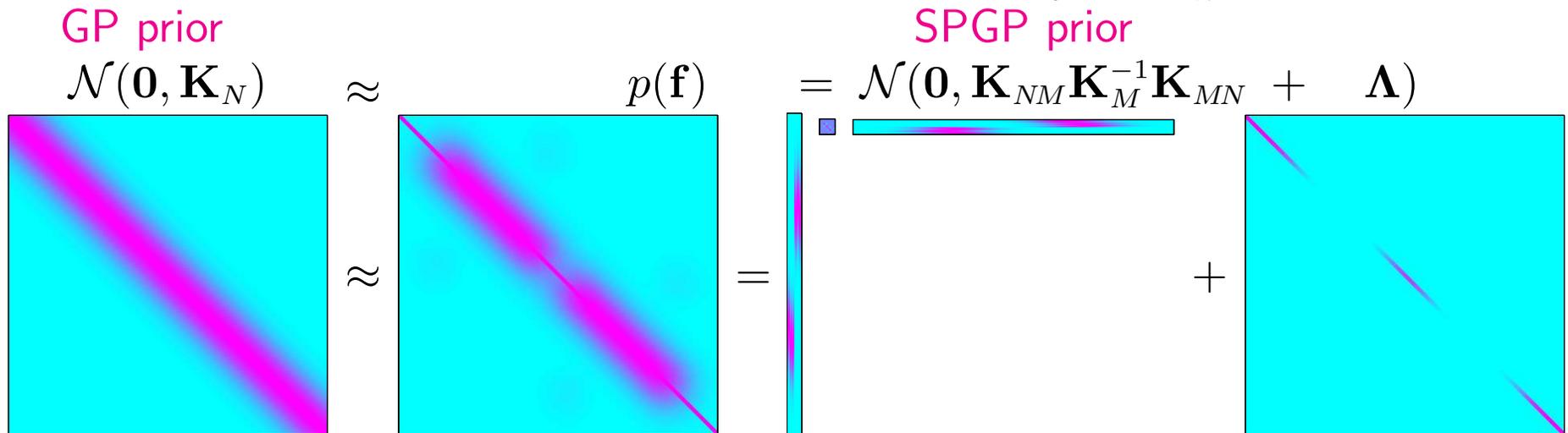
demo\_ep\_2d

demo\_gpr

# Sparse Approximations: Speeding up GP learning

(Snelson and Ghahramani, 2006a, 2006b; Naish-Guzman and Holden 2008)

We can approximate GP through  $M < N$  inducing points  $\bar{\mathbf{f}}$  to obtain this Sparse Pseudo-input Gaussian process (SPGP) prior:  $p(\mathbf{f}) = \int d\bar{\mathbf{f}} \prod_n p(f_n|\bar{\mathbf{f}}) p(\bar{\mathbf{f}})$



- SPGP covariance inverted in  $\mathcal{O}(M^2N) \ll \mathcal{O}(N^3) \Rightarrow$  much faster
- SPGP = GP with non-stationary covariance parameterized by  $\bar{\mathbf{X}}$
- Given data  $\{\mathbf{X}, \mathbf{y}\}$  with noise  $\sigma^2$ , predictive mean and variance can be computed in  $\mathcal{O}(M)$  and  $\mathcal{O}(M^2)$  per test case respectively

Builds on a large lit on sparse GPs (see Quiñonero Candela and Rasmussen, 2006).

# Some Comparisons

Table 1: Test errors and predictive accuracy (smaller is better) for the GP classifier, the support vector machine, the informative vector machine, and the sparse pseudo-input GP classifier.

Data set			GPC		SVM		IVM			SPGPC		
name	train:test	dim	err	nlp	err	#sv	err	nlp	M	err	nlp	M
<i>synth</i>	250:1000	2	0.097	0.227	0.098	98	0.096	0.235	150	<b>0.087</b>	0.234	4
<i>crabs</i>	80:120	5	0.039	0.096	0.168	67	0.066	0.134	60	<b>0.043</b>	0.105	10
<i>banana</i>	400:4900	2	0.105	0.237	0.106	151	<b>0.105</b>	0.242	200	0.107	0.261	20
<i>breast-cancer</i>	200:77	9	0.288	0.558	<b>0.277</b>	122	0.307	0.691	120	0.281	0.557	2
<i>diabetes</i>	468:300	8	0.231	0.475	<b>0.226</b>	271	0.230	0.486	400	0.230	0.485	2
<i>flare-solar</i>	666:400	9	0.346	0.570	<b>0.331</b>	556	0.340	0.628	550	0.338	0.569	3
<i>german</i>	700:300	20	0.230	0.482	0.247	461	0.290	0.658	450	<b>0.236</b>	0.491	4
<i>heart</i>	170:100	13	0.178	0.423	<b>0.166</b>	92	0.203	0.455	120	0.172	0.414	2
<i>image</i>	1300:1010	18	0.027	0.078	0.040	462	<b>0.028</b>	0.082	400	0.031	0.087	200
<i>ringnorm</i>	400:7000	20	0.016	0.071	0.016	157	0.016	0.101	100	<b>0.014</b>	0.089	2
<i>splice</i>	1000:2175	60	0.115	0.281	<b>0.102</b>	698	0.225	0.403	700	0.126	0.306	200
<i>thyroid</i>	140:75	5	0.043	0.093	0.056	61	0.041	0.120	40	<b>0.037</b>	0.128	6
<i>titanic</i>	150:2051	3	0.221	0.514	<b>0.223</b>	118	0.242	0.578	100	0.231	0.520	2
<i>twonorm</i>	400:7000	20	0.031	0.085	0.027	220	0.031	0.085	300	<b>0.026</b>	0.086	2
<i>waveform</i>	400:4600	21	0.100	0.229	0.107	148	0.100	0.232	250	<b>0.099</b>	0.228	10

From (Naish-Guzman and Holden, 2008), using exactly same kernels.

# Feature Selection

**Example:** classification

$$\begin{aligned} \text{input } \mathbf{x} &= (x_1, \dots, x_D) \in \mathbb{R}^D \\ \text{output } y &\in \{+1, -1\} \end{aligned}$$

$2^D$  possible subsets of relevant input features.

One approach, consider all models  $m \in \{0, 1\}^D$  and find

$$\hat{m} = \operatorname{argmax}_m p(\mathcal{D}|m)$$

**Problems:** intractable, overfitting, we should really average

# Feature Selection

- Why are we doing feature selection?
- What does it cost us to keep all the features?
- Usual answer (overfitting) does not apply to fully Bayesian methods, since they don't involve any fitting.
- We should only do feature selection if there is a cost associated with measuring features or predicting with many features.

Note: Radford Neal won the NIPS feature selection competition using Bayesian methods that used 100% of the features.

# Feature Selection using ARD in GPs

**Problem:** Often there are *many* possible inputs that might be relevant to predicting a particular output. We need algorithms that automatically decide which inputs are relevant.

## Automatic Relevance Determination:

Consider this covariance function:

$$\mathbf{K}_{nn'} = v \exp \left[ -\frac{1}{2} \sum_{d=1}^D \left( \frac{x_n^{(d)} - x_{n'}^{(d)}}{r_d} \right)^2 \right]$$

The parameter  $r_d$  is the **length scale of the function along input dimension  $d$** .

As  $r_d \rightarrow \infty$  the function  $f$  varies less and less as a function of  $x^{(d)}$ , that is, the  $d$ th dimension becomes *irrelevant*.

Given data, by learning the lengthscales  $(r_1, \dots, r_D)$  it is possible to do automatic feature selection.

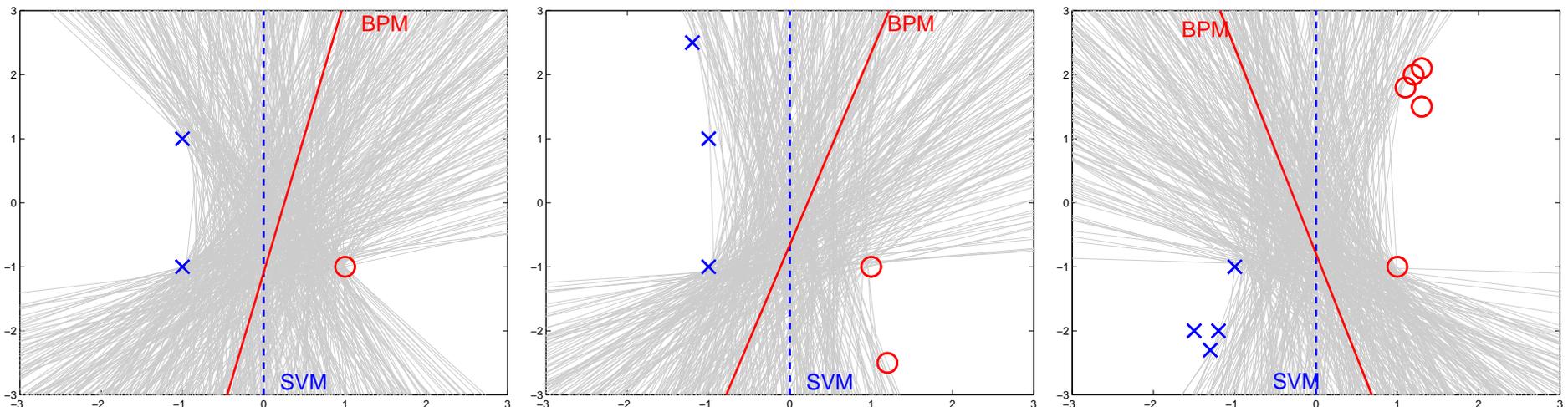
# Bayesian Discriminative Modeling

Terminology for classification with inputs  $\mathbf{x}$  and classes  $y$ :

- **Generative Model:** models prior  $p(y)$  and class-conditional density  $p(\mathbf{x}|y)$
- **Discriminative Model:** directly models the conditional distribution  $p(y|\mathbf{x})$  or the class boundary e.g.  $\{\mathbf{x} : p(y = +1|\mathbf{x}) = 0.5\}$

Myth: Bayesian Methods = Generative Models

For example, it is possible to define Bayesian kernel classifiers (i.e. Gaussian processes) analogous to support vector machines (SVMs).



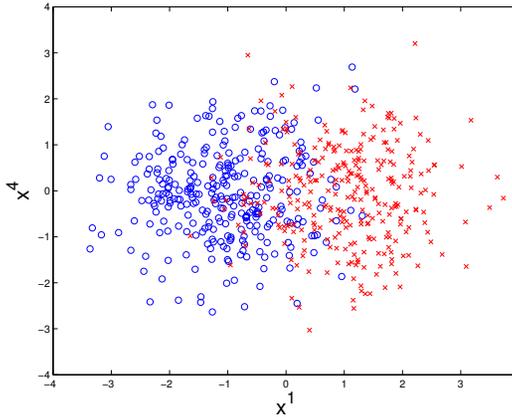
(figure adapted from Minka, 2001)



# Appendix

# An example of ARD for classification

**Data set:** 6-dimensional data set with three *relevant* features and three *irrelevant* features. For each data point  $\vec{x}_i$ , the relevant features depend on its class label:  $x_i^1, x_i^2, x_i^3 \sim \mathcal{N}(y_i, 1)$ , while the irrelevant features do not:  $x_i^4, x_i^5, x_i^6 \sim \mathcal{N}(0, 1)$ .



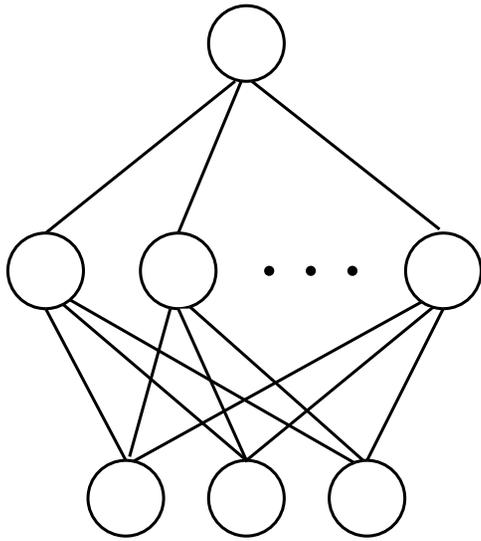
**Result:**  $r_4, r_5, r_6 \rightarrow \infty$  improving the likelihood and classification error rates, compared to a single-lengthscale model.

Methods	single lengthscale	multiple lengthscales
$\log p(\mathbf{y} \mathbf{X}, \boldsymbol{\theta})$	-55.4480	-35.4119
Error rates	0.0600	0.0400

Example from (Kim and Ghahramani, 2004)

More on ARD and feature selection with thousands of inputs: (Qi et al, 2004).

# Feature Selection: Automatic Relevance Determination



## Bayesian neural network

Data:  $\mathcal{D} = \{(\mathbf{x}^{(n)}, y^{(n)})\}_{n=1}^N = (X, \mathbf{y})$

Parameters (weights):  $\boldsymbol{\theta} = \{\{w_{ij}\}, \{v_k\}\}$

prior  $p(\boldsymbol{\theta}|\boldsymbol{\alpha})$

posterior  $p(\boldsymbol{\theta}|\boldsymbol{\alpha}, \mathcal{D}) \propto p(\mathbf{y}|X, \boldsymbol{\theta})p(\boldsymbol{\theta}|\boldsymbol{\alpha})$

evidence  $p(\mathbf{y}|X, \boldsymbol{\alpha}) = \int p(\mathbf{y}|X, \boldsymbol{\theta})p(\boldsymbol{\theta}|\boldsymbol{\alpha}) d\boldsymbol{\theta}$

prediction  $p(y'|\mathcal{D}, \mathbf{x}', \boldsymbol{\alpha}) = \int p(y'|\mathbf{x}', \boldsymbol{\theta})p(\boldsymbol{\theta}|\mathcal{D}, \boldsymbol{\alpha}) d\boldsymbol{\theta}$

## Automatic Relevance Determination (ARD):

Let the weights from feature  $x_d$  have variance  $\alpha_d^{-1}$ :  $p(w_{dj}|\alpha_d) = \mathcal{N}(0, \alpha_d^{-1})$

Let's think about this:

$\alpha_d \rightarrow \infty$	variance $\rightarrow 0$	weights $\rightarrow 0$	(irrelevant)
$\alpha_d \ll \infty$	finite variance	weight can vary	(relevant)

**ARD:** optimize  $\hat{\boldsymbol{\alpha}} = \underset{\boldsymbol{\alpha}}{\operatorname{argmax}} p(\mathbf{y}|X, \boldsymbol{\alpha})$ .

During optimization some  $\alpha_d$  will go to  $\infty$ , so the model will discover irrelevant inputs.

# Sparse GP overview

This work contains 2 key ideas:

1. A new sparse Gaussian process approximation based on a small set of  $M$  'pseudo-inputs' ( $M \ll N$ ). This reduces computational complexity to  $\mathcal{O}(M^2N)$
2. A gradient based learning procedure for finding the pseudo-inputs and hyperparameters of the Gaussian process, in one joint optimization

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- More information and code at: <http://www.gaussianprocess.org/>