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Equivalent Kernel

Abstract—This document contains theory involved in curve fitting.

1 Objective

The objective is to use an equivalent kernel on a noisy data.

2 Generate Dataset

Create a sinusoidal function of the form

$$y = A \sin 2\pi x + n(t)$$
 (2.0.1)

n(t) is the random noise that is included in the training set. This set consists of N samples of input data i.e. x expressed as shown below

$$x = (x_1, x_2, ..., x_N)^T$$
 (2.0.2)

which give the corresponding values of y denoted as

$$y = (y_1, y_2, ..., y_N)^T$$
 (2.0.3)

The corresponding values of y are generated from the Eq (2.0.1). The first term $A \sin 2\pi x$ is computed directly and then random noise samples having a normal (Gaussian) distribution are added in order to get the corresponding values of y.

#Generate the sine curve

noise = 0.5

A = 2

Noisy training data

X train = np.arange(-3, 4, 1).reshape(-1, 1)

Y_train = A*np.sin(2*np.pi*X_train) + noise * np.random.randn(*X_train.shape)

The generated input matrix would look like

$$\mathbf{F} = \begin{pmatrix} 1 & x_0 & x_0^2 & \dots & x_0^{N-1} \\ 1 & x_1 & x_1^2 & \dots & x_1^{N-1} \\ 1 & x_2 & x_2^2 & \dots & x_2^{N-1} \\ \vdots & \vdots & & \vdots \\ 1 & \dots & \dots & x_N^{N-1} \end{pmatrix}$$
(2.0.4)

3 POLYNOMIAL CURVE FITTING

The goal is to find the best line that fits into the pattern of the training data shown in the graph. We shall fit the data using a polynomial function of the form,

$$y(w, x) = \sum_{j=0}^{M} w_j x^j$$
 (3.0.1)

(3.0.2)

M is the order of the polynomial The polynomial coefficient are collectively denoted by the vector \mathbf{w} . The proposed vector \mathbf{w} of the model referring to Eq (2.0.4) is given by

$$\hat{\mathbf{w}} = \left(\mathbf{F}^T \mathbf{F}\right)^{-1} \mathbf{F}^T y \tag{3.0.3}$$

4 Equivalent Kernel

For Bayesian treatment of linear regression, we introduce a prior probability distribution over the model parameters **w**.

The corresponding conjugate prior is therefore given by a Gaussian distribution of the form

$$p(\mathbf{w}) = N(\mathbf{w}|\mathbf{m_0}, \mathbf{S_0}) \tag{4.0.1}$$

having mean \mathbf{m}_0 and covariance \mathbf{S}_0 .

We now compute the posterior distribution, which is proportional to the product of the likelihood function and the prior which takes the form

$$p(\mathbf{w}|t) = N(\mathbf{w}|\mathbf{m}_{\mathbf{N}}, \mathbf{S}_{\mathbf{N}}) \tag{4.0.2}$$

Since the prior is Gaussian, so would be the posterior. In Eq (4.0.2),

$$\mathbf{m}_N = \mathbf{S}_N \left(\mathbf{S}_0^{-1} \mathbf{m}_0 + \beta f^T t \right) \tag{4.0.3}$$

$$\mathbf{S}_{N}^{-1} = \mathbf{S}_{0}^{-1} + \beta f^{T} f \tag{4.0.4}$$

Specifically, we consider a zero - mean isotropic Gaussian governed by a single precision parameter α so that

$$p(\mathbf{w}|\alpha) = N(\mathbf{w}|0, \alpha^{-1}I)$$
 (4.0.5)

Then the corresponding posterior distribution over where w is then given as

$$\mathbf{m}_N = \beta \mathbf{S}_N f^T t \tag{4.0.6}$$

$$\mathbf{S}_N^{-1} = \alpha I + \beta f^T f \tag{4.0.7}$$

The predictive mean can be written in the form

$$y(\mathbf{x}, \mathbf{m}_N) = \mathbf{m}_N f(x)$$

$$= \beta f(x)^T S_N f^T t$$

$$= \sum_{n=1}^N \beta f(x)^T S_N f(\mathbf{x}_n) t_n \quad (4.0.8)$$

The mean of the predictive distribution at a point \mathbf{x} is given by a linear combination of the training set target variables t_n , so

$$y(\mathbf{x}, \mathbf{m}_N) = \sum_{n=1}^{N} k(\mathbf{x}, \mathbf{x}') t_n$$
 (4.0.9)

where

$$k(\mathbf{x}, \mathbf{x}_n) = \beta f(x)^T S_N f(\mathbf{x}')$$
 (4.0.10)

Eq (4.0.10) is known as the smoother matrix of the equivalent kernel.

Further insight into the role of the equivalent kernel can be obtained by considering the covariance between $y(\mathbf{x})$ and $y(\mathbf{x}')$ given by

$$cov[y(\mathbf{x}), y(\mathbf{x}')] = cov[f(\mathbf{x})^T \mathbf{w}, \mathbf{w}^T f(\mathbf{x}')] \quad (4.0.11)$$
$$= f(\mathbf{x})^T S_N f(\mathbf{x}') \quad (4.0.12)$$
$$= \beta^{-1} k(\mathbf{x}, \mathbf{x}') \quad (4.0.13)$$

we define a localized kernel directly and use this to make predictions for new input vectors x.

This leads to practical frameworks for regression known as Gaussian Processes.

An effective kernel should satisfy

$$\sum_{n=1}^{N} k(\mathbf{x}, \mathbf{x}_n) = 1 \tag{4.0.14}$$

for all values of x.

Eq (4.0.10) satisfies the general equation of kernel, that can be expressed in the form an inner product with respect to a vector $f(\mathbf{x})$ of non linear functions given by

$$k(\mathbf{x}, \mathbf{z}) = f(\mathbf{x})^T f(\mathbf{z}) \tag{4.0.15}$$

$$f(\mathbf{x}) = \beta^{\frac{1}{2}} S_N^{\frac{1}{2}} f(\mathbf{x})$$
 (4.0.16)

5 Implementation using numPy

Here, we will use the squared exponential kernel, also known as Gaussian kernel, given by

$$k(\mathbf{x}_i, \mathbf{x}_j) = \sigma_f^2 \exp{-\frac{1}{2l^2}(\mathbf{x}_i - \mathbf{x}_j)^T(\mathbf{x}_i - \mathbf{x}_j)}$$
 (5.0.1)

The length *l* controls the smoothness of the function and σ_f the vertical variation

We define the kernel Eq (5.0.1),

```
#Kernel
def kernel(X1, X2, l=1.0, sigma f=1.0):
    sqdist = np.sum(X1**2, 1).reshape(-1, 1) +
        np.sum(X2**2, 1) - 2 * np.dot(X1, X2.T)
    return sigma f**2 * np.exp(-0.5 / 1**2 *
        sqdist)
```

We use a function for plotting the samples

```
def plot gp(mu, cov, X, X train=None,
    Y train=None, samples=[]):
   X = X.ravel()
   mu = mu.ravel()
   uncertainty = 1.96 * np.sqrt(np.diag(cov))
   plt.fill between(X, mu + uncertainty, mu -
       uncertainty, alpha=0.1)
   plt.plot(X, mu, label='Mean')
   for i, sample in enumerate(samples):
       plt.plot(X, sample, lw=1, ls='--', label=
           f'Sample \{i+1\}')
   if X train is not None:
       plt.plot(X train, Y train, 'rx')
   plt.legend()
```

The following code draws three random samples and plots it together with the zero mean.

```
import numpy as np
import matplotlib.pyplot as plt
X = \text{np.arange}(-5, 5, 0.2).\text{reshape}(-1, 1)
mu = np.zeros(X.shape)
cov = kernel(X, X)
# Draw three samples from the prior
```

samples = np.random.multivariate_normal(mu. ravel(), cov, 3)

Plot GP mean, confidence interval and samples plot_gp(mu, cov, X, samples=samples)

The Plot would look as below shown in Fig 0

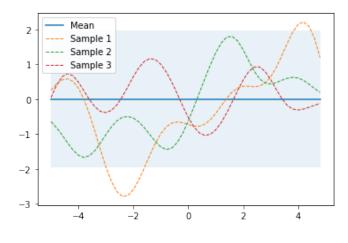


Fig. 0

We define the posterior as

from numpy.linalg import inv

def posterior(X_s, X_train, Y_train, l=1.0,
 sigma_f=1.0, sigma_y=1e-8):
 K = kernel(X_train, X_train, l, sigma_f) +
 sigma_y**2 * np.eye(len(X_train))
 K_s = kernel(X_train, X_s, l, sigma_f)
 K_ss = kernel(X_s, X_s, l, sigma_f) + 1e
 -8 * np.eye(len(X_s))
 K_inv = inv(K)

Equation (7)
 mu_s = K_s.T.dot(K_inv).dot(Y_train)

Equation (8)
 cov_s = K_ss - K_s.T.dot(K_inv).dot(K_s
)

Now, the prediction from noisy training data

return mu s, cov s

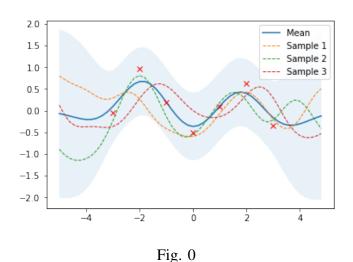
Y_train = A*np.sin(2*np.pi*X_train) + noise *
np.random.randn(*X train.shape)

Compute mean and covariance of the posterior distribution

samples = np.random.multivariate_normal(mu_s. ravel(), cov_s, 3)

plot_gp(mu_s, cov_s, X, X_train=X_train, Y_train=Y_train, samples=samples)

Th plot would look like in Fig 0



Python code:

https://github.com/Hrithikraj2/EE4015_IDP/blob/main/Assignment 6/Assignment 6.ipynb