GP lab Milda

Milda Poceviciute
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Question 1

Part 1

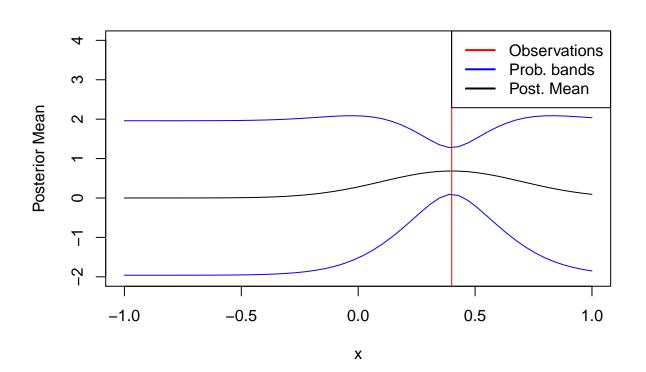
The code below is the implementation of the Gaussian process regression model:

```
y = f(x) + \epsilon with \epsilon \sim N(0, \sigma_n^2), f \sim GP(0, k(x, x'))
# Setting up the kernel
SquaredExpKernel <- function(x1,x2,sigmaF=1,l=3){</pre>
    n1 <- length(x1)
    n2 \leftarrow length(x2)
    K <- matrix(NA,n1,n2)</pre>
    for (i in 1:n2){
        K[,i] \leftarrow sigmaF^2*exp(-0.5*((x1-x2[i])/1)^2)
    }
    return(K)
}
posteriorGP <- function(X, y, XStar, hyperParam, sigmaNoise) {</pre>
    k_star <- SquaredExpKernel(X, XStar, sigmaF = hyperParam$sigmaF, 1 = hyperParam$1)
    K <- SquaredExpKernel(X, X, sigmaF = hyperParam$sigmaF, 1 = hyperParam$1)</pre>
    noise <- diag(nrow(K)) * sigmaNoise</pre>
    L <- t(chol(K + noise))
    # Predictive mean
    alpha <- solve(L) %*% y
    posterior_mean <- t(k_star) %*% alpha</pre>
    # Predictive variance
    v <- solve(L) %*% k star
    kk <- SquaredExpKernel(XStar, XStar, sigmaF = hyperParam$sigmaF, 1 = hyperParam$1)
    posterior_var <- kk - t(v) %*% v
    return(list(posterior_mean = posterior_mean, posterior_var = posterior_var))
}
```

Part 2

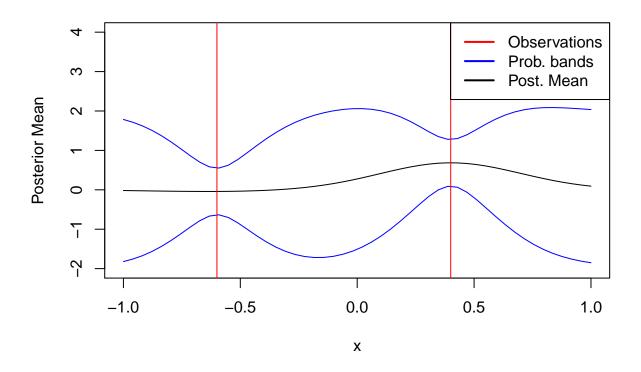
In this part we assume that prior hyper-parameters are $\sigma_f = 1$, $\ell = 0.3$, and we update the posterior with observation (x, y) = (0.4, 0.719). Here σ_n is assumed to be 0.1. The plot below shows the observation, the posterior mean (the black line), and the 95% probability bands (point-wise) plotted over a grid [-1, 1]. The probability bands are quite wide for most of the x grid points, except from the location of the observed point (x, y). This observation decreases the uncertainty and bands become narrower.

```
# Information for updating the prior
X observations \leftarrow c(0.4)
y_observations <- c(0.719)
hyperParam2 <- list(sigmaF = 1, 1 = 0.3)
# Use the data set and new given point to compute the posterior over the grid [-1,1]
xGrid \leftarrow seq(-1,1,length=50)
posteriorP2 <- posteriorGP(X = X_observations, y = y_observations, XStar = xGrid, hyperParam = hyperPar
# find probability bounds
error <- qnorm(0.975)*sqrt(diag(posteriorP2$posterior_var))</pre>
upp_band <- as.vector(posteriorP2$posterior_mean) + as.vector(error)</pre>
low_band <- as.vector(posteriorP2$posterior_mean) - as.vector(error)</pre>
# Plot the posterior means over a grid
plot(x=xGrid,y=posteriorP2$posterior_mean, type="1",ylim=c(-2,4),xlab="x", ylab="Posterior Mean")
# Add 95% probability bounds (check bayesian course, we must have doen it there)
lines(x=xGrid,y=upp_band, type="1",col="blue")
lines(x=xGrid,y=low_band, type="l",col="blue")
abline(v=0.4,col="red")
legend("topright",c("Observations","Prob. bands","Post. Mean"),
        col = c("red","blue","black"), lty = c(1, 1, 1), lwd=c(2,2,2))
```



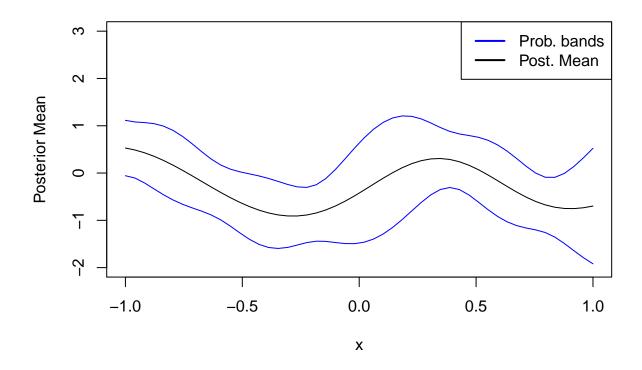
The posterior is updated with one more observation (x, y) = (-0.6, -0.044). The plot below shows again the observations, the posterior mean(the black line), and the 95% probability bands (point-wise) over the grid [-1,1]. This time the uncertainty is smaller for the two observed x values: the probability bands are narrower there.

```
# Update the information
X_observations[2] <- -0.6</pre>
y_observations[2] <- -0.044
posteriorP3 <- posteriorGP(X = X_observations, y = y_observations, XStar = xGrid, hyperParam = hyperPar
# find probability bounds
error3 <- qnorm(0.975)*sqrt(diag(posteriorP3$posterior_var))</pre>
upp_band3 <- as.vector(posteriorP3$posterior_mean) + as.vector(error3)</pre>
low_band3 <- as.vector(posteriorP3$posterior_mean) - as.vector(error3)</pre>
# Plot the posterior means over a grid
plot(x=xGrid,y=posteriorP3$posterior_mean, type="1",ylim=c(-2,4),xlab="x", ylab="Posterior Mean")
# Add 95% probability bounds (check bayesian course, we must have doen it there)
lines(x=xGrid,y=upp_band3, type="l",col="blue")
lines(x=xGrid,y=low_band3, type="l",col="blue")
abline(v=0.4,col="red")
abline(v=-0.6,col="red")
legend("topright",c("Observations","Prob. bands","Post. Mean"),
        col = c("red","blue","black"), lty = c(1, 1, 1), lwd=c(2,2,2))
```

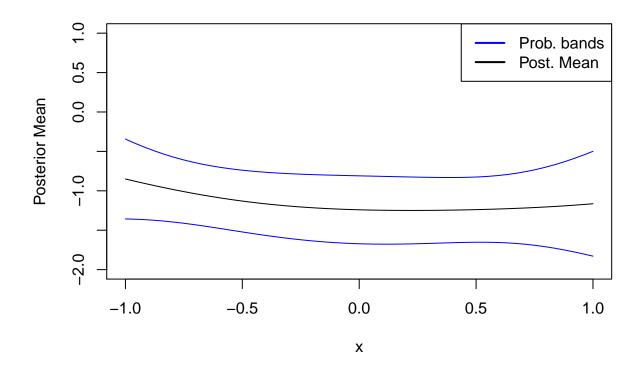


Part 4

The plot shows the posterior mean and the 95% probability bands after having 5 observations. The uncertainty has decreased a lot in comparison to parts 2 and 3.



The plot below shows the posterior mean and the 95% probability bands after having the 5 observations as in part 4, but changing the hyper-parameters to $\sigma_f = 1$, $\ell = 1$. The higher the ℓ value, the smoother the posterior mean becomes. In the kernel the distance between the points is divided by ℓ . This means that the higher the value of this hyper-parameter is, the kernel values converge to the σ_f : the distance between the observations becomes less and less important. Hence, the posterior mean becomes more smooth.



Question 2

Part 1

```
dataQ2 <- read.csv("TempTullinge.csv", header=TRUE, sep=";")
time <- 1:2190
day <- 1:365
Stime <- seq(1,2190,by=5)
Sday <- seq(1,365,by=5)
x1 <- c(1,3,4)
x2 <- c(2,3,4)</pre>
```

In this question, the exponential kernel function is defined:

```
# Function that returns kernel class object
Matern32 <- function(sigmaf = 1, ell = 1)
{
    rval <- function(x1,x2){
        n1 <- length(x1)
        n2 <- length(x2)
        K <- matrix(NA,n1,n2)
        for (i in 1:n2){
            K[,i] <- sigmaf^2*exp(-0.5*( (x1-x2[i])/ell)^2 )
        }
        return(K)</pre>
```

```
class(rval) <- "kernel"</pre>
  return(rval)
}
MaternFunc = Matern32(sigmaf = 1, ell = 2)
It is used to compute the covariance, firstly between (1,2), and afterwards between points x=(1,3,4) and
x^* = (2, 3, 4). The results are printed:
#evaluate it in the point x = 1; x' = 2
cat("The kernel between points (1,2) is :", MaternFunc(1,2))
## The kernel between points (1,2) is: 0.8824969
cat("\n")
cat("\n")
# use the kernelMatrix function to compute the covariance matrix K(X;X*)
covM <- kernelMatrix(MaternFunc, x1,x2)</pre>
cat("The kernel between points (1,3,4) and (2,3,4) is:")
## The kernel between points (1,3,4) and (2,3,4) is:
cat("\n")
covM
## An object of class "kernelMatrix"
                        [,2]
              [,1]
## [1,] 0.8824969 0.6065307 0.3246525
```

The following model is used for obtaining the posterior mean:

[2,] 0.8824969 1.0000000 0.8824969 ## [3,] 0.6065307 0.8824969 1.0000000

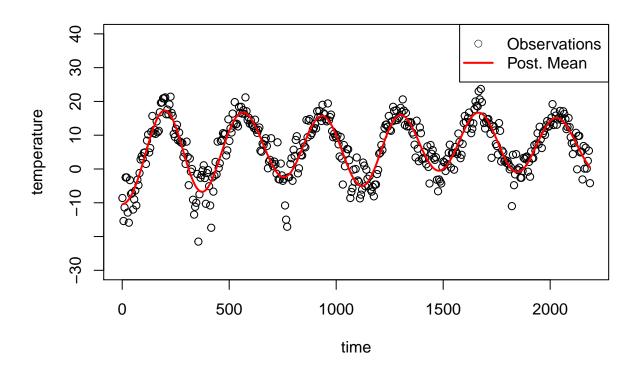
```
temp = f(time) + \epsilon \text{ with } \epsilon \sim N(0, \sigma_n^2), f \sim GP(0, k(time, time'))
```

```
# Subset data
temp <- dataQ2$temp[Stime]
# Find the noise value
polyFit <- lm(temp ~ Stime + Stime^2)
sigmaNoise = sd(polyFit$residuals)
MaternFunc2 = Matern32(sigmaf = 20, ell = 0.2)

# Fit the GP with built in Square expontial kernel (called rbfdot in kernlab)
GPfit <- gausspr(Stime, temp, kernel = MaternFunc2, var = sigmaNoise^2)
postMean2 <- predict(GPfit,Stime)</pre>
```

The plot below shows the observations plotted with a fitted posterior mean at each data point. The model fits the data quite well as it predicts the underlying trends (periodicity).

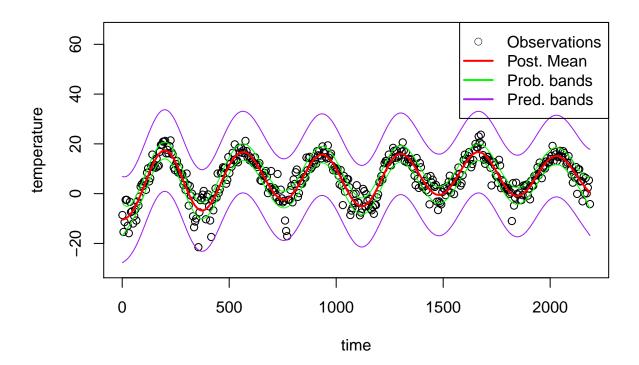
```
plot(Stime, temp, ylim=c(-30,40),xlab="time",ylab="temperature")
lines(Stime,postMean2, col="red", lwd = 2)
```



```
x<-scale(Stime)
xs<-scale(Stime) # XStar
n <- length(x)
SEkernel <- MaternFunc2
Kss <- kernelMatrix(kernel = SEkernel, x = xs, y = xs)
Kxx <- kernelMatrix(kernel = SEkernel, x = x, y = x)
Kxs <- kernelMatrix(kernel = SEkernel, x = x, y = xs)
Covf = Kss-t(Kxs)%*%solve(Kxx + sigmaNoise^2*diag(n), Kxs) # Covariance matrix of fStar</pre>
```

From the plot below, we see that the probability and prediction bands fit the data quite well. Though, the prediction bands are quite wide.

```
plot(Stime, temp, ylim=c(-30,65),xlab="time",ylab="temperature")
lines(Stime,postMean2, col="red", lwd = 2)
# Probability intervals for fStar
lines(Stime, postMean2 - 1.96*sqrt(diag(Covf)), col = "green")
lines(Stime, postMean2 + 1.96*sqrt(diag(Covf)), col = "green")
# Prediction intervals for yStar
lines(Stime, postMean2 - 1.96*sqrt((diag(Covf) + sigmaNoise^2)), col = "purple")
lines(Stime, postMean2 + 1.96*sqrt((diag(Covf) + sigmaNoise^2)), col = "purple")
```



Part 4

The following model is used for obtaining the posterior mean:

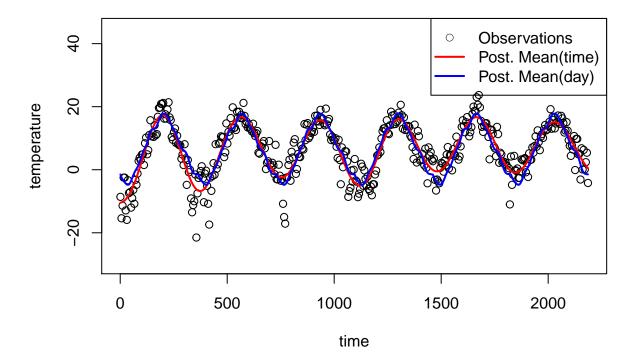
```
temp = f(day) + \epsilon \text{ with } \epsilon \sim N(0, \sigma_n^2), f \sim GP(0, k(day, day'))
```

```
# Find the noise value
polyFit4 <- lm(temp ~ rep.int(Sday,6) )
sigmaNoise4 = sd(polyFit4$residuals)

# Fit the GP with built in Square expontial kernel (called rbfdot in kernlab)
GPfit4 <- gausspr(rep.int(Sday,6), temp, kernel = MaternFunc2, var = sigmaNoise4^2)
postMean4 <- predict(GPfit4,rep.int(Sday,6))</pre>
```

From the plot, we see that both models performed quite well and the posterior means follow closely the seasonality of the temperatures. The *Time* model was fitting to the all observed information, hence it shows the overall trend of the temperature fluctuations over the whole time period. The *Day* model captures the seasonality during the year, as the variable 1, 2, ..., 365 was repeated 6 times and fitted with the temperatures. It fits exactly the same curve 6 times.

```
plot(Stime, temp,xlab="time",ylab="temperature",ylim=c(-30,45))
lines(Stime,postMean2, col="red", lwd = 2)
lines(Stime,postMean4, col="blue", lwd = 2)
```



Part 5

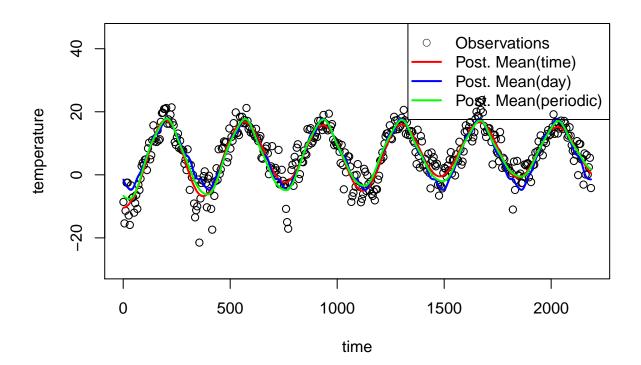
The GP with a generalization of the periodic kernel is used in this part to model the temperature against the time. The periodic kernel is given by:

$$k(x, x') = \sigma_f^2 \exp\left\{-\frac{2\sin^2(pi|x - x'|)/d}{\ell_1^2}\right\} \exp\left\{-\frac{|x - x'|^2}{2\ell_2^2}\right\}$$

```
# Function that returns kernel class object
Matern5 <- function(sigmaf, ell1, ell2, d){
    val <- function(x1, x2){
        n1 <- length(x1)
        n2 <- length(x2)
        K <- matrix(NA,n1,n2)
        for (i in 1:n2){
            K[,i] <- sigmaf^2*exp(-2*sin(pi*abs(x1-x2[i])/d)^2/ell1^2)*exp(-0.5*abs(x1-x2[i])^2/ell2^2)
        }
        return(K)
    }
    class(val) <- "kernel"
    return(val)
}</pre>
```

```
MaternFunc5 <- Matern5(20, 1, 10, 365/sd(time))
GPfit5 <- gausspr(Stime, temp, kernel = MaternFunc5, var = sigmaNoise^2)
postMean5 <- predict(GPfit5,Stime)</pre>
```

From the plot, we see that the posterior mean from *Time* model predicts the underlying trends over all time period, the *Day* model is good at capturing the yearly seasonality. The *Periodic* model of this part seems to be performing quite well too. It seems the model with periodic kernel captures the overall trend as the model with the squared kernel with variable *Time*. But also it shows the average year seasonality as the model with squared kernel and variable *Day*. Since periodic kernel has one more length scale which controls the correlation between the same day in different years, this result looks reasonable.



Question 3

```
data[,5] <- as.factor(data[,5])</pre>
head(data)
     varWave skewWave kurtWave entropyWave fraud
## 1 3.62160 8.6661 -2.8073
                                  -0.44699
## 2 4.54590 8.1674 -2.4586
                                  -1.46210
                                               0
## 3 3.86600 -2.6383 1.9242
                                   0.10645
                                               0
## 4 3.45660 9.5228 -4.0112
                                  -3.59440
                                               0
## 5 0.32924 -4.4552 4.5718
                                  -0.98880
## 6 4.36840 9.6718 -3.9606
                                  -3.16250
                                               0
set.seed(111);
SelectTraining <- sample(1:dim(data)[1], size = 1000,
                                        replace = FALSE)
training <- data[SelectTraining,]</pre>
testing <- data[-SelectTraining,]</pre>
accuracy <- function (conf_mat){(conf_mat[1,1] + conf_mat[2,2]) / sum(conf_mat)}</pre>
```

The package *kenrlab* is used to fit the Gaussian Process Classification on Fraud data-set.

```
GPfitFraud <- gausspr(fraud ~ varWave + skewWave, data=training)</pre>
```

Using automatic sigma estimation (sigest) for RBF or laplace kernel

```
# class probabilities
probPreds <- predict(GPfitFraud, training, type="probabilities")
x1 <- seq(min(training[,1]),max(training[,1]),length=100)
x2 <- seq(min(training[,2]),max(training[,2]),length=100)
gridPoints <- meshgrid(x1, x2)
gridPoints <- cbind(c(gridPoints$x), c(gridPoints$y))

gridPoints <- data.frame(gridPoints)
names(gridPoints) <- names(training)[1:2]
probPreds <- predict(GPfitFraud, gridPoints, type="probabilities")</pre>
```

The model gives the corresponding confusion matrix on the training data:

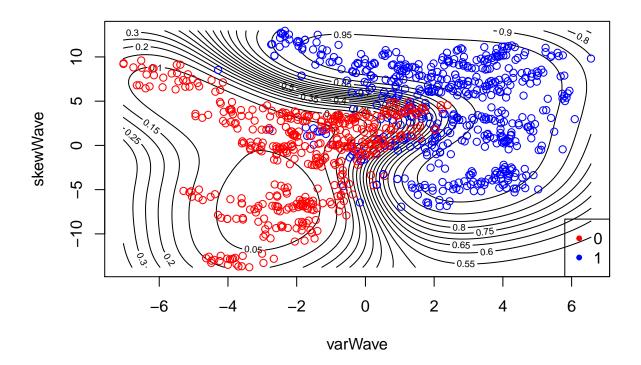
```
# predict on the testing set
pred1 <- predict(GPfitFraud,training)
conM1 <- table("Prediction"=pred1, "True"=training$fraud) # confusion matrix
conM1</pre>
```

```
## True
## Prediction 0 1
## 0 512 24
## 1 44 420
```

Hence, the accuracy is 0.932. From the plot below, we see that classifier works quite well and there are two contour peaks that concentrate the most of the points from the different class. However, the overlap region is quite big, and that explains the lower accuracy rate of the classifier.

```
points(training[training[,5]==0,1],training[training[,5]==0,2],col="blue")
points(training[training[,5]==1,1],training[training[,5]==1,2],col="red")
legend("bottomright", legend=c("0","1"), pch=c(20,20), col=c("red","blue"))
```

Contour of Prediction Probabilities



Part 2

The confusion matrix on the testing data is:

```
# predict on the testing set
predT <- predict(GPfitFraud,testing)
conM2 <- table("Prediction"=predT, "True"=testing$fraud) # confusion matrix
conM2

## True
## Prediction 0 1
## 0 191 9
## 1 15 157</pre>
```

The accuracy is 0.9354839, which is pretty close to the accuracy rate obtained on the training data. This is a good indication that the classification works well, and it is not overfitted.

Part 3

In this part all available information in the data-set is used to classify the fraud. The resulting confusion matrix on the test data is:

```
GPfitFraud3 <- gausspr(fraud ~ ., data=training)

## Using automatic sigma estimation (sigest) for RBF or laplace kernel

# predict on the testing set

pred3 <- predict(GPfitFraud3,testing)
conM3 <- table("Prediction"=pred3, "True"= testing$fraud) # confusion matrix
conM3

## True

## Prediction 0 1

## 0 205 0

## 1 1 166</pre>
```

The accuracy is 0.9973118, which is higher than in the previous parts, and very close to 1. It is reasonable that the classification model improved once we included all the available information.

Appendix

```
knitr::opts_chunk$set(echo = TRUE)
library(kernlab)
library(AtmRay)
set.seed(12345)
# Setting up the kernel
SquaredExpKernel <- function(x1,x2,sigmaF=1,l=3){</pre>
    n1 \leftarrow length(x1)
    n2 \leftarrow length(x2)
    K <- matrix(NA,n1,n2)</pre>
    for (i in 1:n2){
        K[,i] \leftarrow sigmaF^2*exp(-0.5*((x1-x2[i])/1)^2)
    }
    return(K)
}
posteriorGP <- function(X, y, XStar, hyperParam, sigmaNoise) {</pre>
    k_star <- SquaredExpKernel(X, XStar, sigmaF = hyperParam$sigmaF, 1 = hyperParam$1)</pre>
    K <- SquaredExpKernel(X, X, sigmaF = hyperParam$sigmaF, 1 = hyperParam$1)</pre>
    noise <- diag(nrow(K)) * sigmaNoise</pre>
    L <- t(chol(K + noise))</pre>
    # Predictive mean
    alpha <- solve(L) %*% y
    posterior_mean <- t(k_star) %*% alpha</pre>
    # Predictive variance
    v <- solve(L) %*% k_star</pre>
    kk <- SquaredExpKernel(XStar, XStar, sigmaF = hyperParam$sigmaF, 1 = hyperParam$1)
    posterior_var <- kk - t(v) %*% v</pre>
    return(list(posterior_mean = posterior_mean, posterior_var = posterior_var))
}
```

```
# Information for updating the prior
X observations \leftarrow c(0.4)
y observations \leftarrow c(0.719)
hyperParam2 <- list(sigmaF = 1, 1 = 0.3)
# Use the data set and new given point to compute the posterior over the grid [-1,1]
xGrid \leftarrow seq(-1,1,length=50)
posteriorP2 <- posteriorGP(X = X_observations, y = y_observations, XStar = xGrid, hyperParam = hyperPar
# find probability bounds
error <- qnorm(0.975)*sqrt(diag(posteriorP2$posterior_var))</pre>
upp_band <- as.vector(posteriorP2$posterior_mean) + as.vector(error)</pre>
low_band <- as.vector(posteriorP2$posterior_mean) - as.vector(error)</pre>
# Plot the posterior means over a grid
plot(x=xGrid,y=posteriorP2$posterior_mean, type="1",ylim=c(-2,4),xlab="x", ylab="Posterior Mean")
# Add 95% probability bounds (check bayesian course, we must have doen it there)
lines(x=xGrid,y=upp_band, type="1",col="blue")
lines(x=xGrid,y=low_band, type="l",col="blue")
abline(v=0.4,col="red")
legend("topright",c("Observations","Prob. bands","Post. Mean"),
        col = c("red","blue","black"), lty = c(1, 1, 1), lwd=c(2,2,2))
# Update the information
X_observations[2] <- -0.6</pre>
y_observations[2] <- -0.044
posteriorP3 <- posteriorGP(X = X_observations, y = y_observations, XStar = xGrid, hyperParam = hyperPar
# find probability bounds
error3 <- qnorm(0.975)*sqrt(diag(posteriorP3$posterior_var))</pre>
upp_band3 <- as.vector(posteriorP3$posterior_mean) + as.vector(error3)</pre>
low_band3 <- as.vector(posteriorP3$posterior_mean) - as.vector(error3)</pre>
# Plot the posterior means over a grid
plot(x=xGrid,y=posteriorP3$posterior_mean, type="1",ylim=c(-2,4),xlab="x", ylab="Posterior Mean")
# Add 95% probability bounds (check bayesian course, we must have doen it there)
lines(x=xGrid,y=upp_band3, type="1",col="blue")
lines(x=xGrid,y=low_band3, type="1",col="blue")
abline(v=0.4,col="red")
abline(v=-0.6,col="red")
legend("topright",c("Observations","Prob. bands","Post. Mean"),
        col = c("red","blue","black"), lty = c(1, 1, 1), lwd=c(2,2,2))
X_observations[1] <- -1</pre>
X_{observations}[3:5] <- c(-0.2,0.4,0.8)
y_observations[3:5] < c(-0.940, 0.719, -0.664)
posteriorP4 <- posteriorGP(X = X_observations, y = y_observations, XStar = xGrid, hyperParam = hyperPar
# find probability bounds
error4 <- qnorm(0.975)*sqrt(diag(posteriorP4$posterior_var))</pre>
```

```
upp_band4 <- as.vector(posteriorP4$posterior_mean) + as.vector(error4)</pre>
low_band4 <- as.vector(posteriorP4$posterior_mean) - as.vector(error4)</pre>
# Plot the posterior means over a grid
plot(x=xGrid,y=posteriorP4$posterior_mean, type="1",ylim=c(-2,3),xlab="x", ylab="Posterior Mean")
# Add 95% probability bounds (check bayesian course, we must have doen it there)
lines(x=xGrid,y=upp_band4, type="1",col="blue")
lines(x=xGrid,y=low band4, type="1",col="blue")
legend("topright",c("Prob. bands","Post. Mean"),
        col = c("blue","black"), lty = c(1, 1), lwd=c(2,2))
hyperParam5 <- list(sigmaF = 1, l = 1)</pre>
posteriorP5 <- posteriorGP(X = X_observations, y = y_observations, XStar = xGrid, hyperParam = hyperPar
# find probability bounds
error5 <- qnorm(0.975)*sqrt(diag(posteriorP5$posterior_var))</pre>
upp_band5 <- as.vector(posteriorP5$posterior_mean) + as.vector(error5)</pre>
low_band5 <- as.vector(posteriorP5$posterior_mean) - as.vector(error5)</pre>
# Plot the posterior means over a grid
plot(x=xGrid,y=posteriorP5$posterior_mean, type="l", ylim = c(-2,1),xlab="x", ylab="Posterior Mean")
# Add 95% probability bounds (check bayesian course, we must have doen it there)
lines(x=xGrid,y=upp_band5, type="1",col="blue")
lines(x=xGrid,y=low_band5, type="l",col="blue")
legend("topright",c("Prob. bands","Post. Mean"),
        col = c("blue", "black"), lty = c(1, 1), lwd=c(2,2))
dataQ2 <- read.csv("TempTullinge.csv", header=TRUE, sep=";")</pre>
time <- 1:2190
day <- 1:365
Stime <- seq(1,2190,by=5)
Sday \leftarrow seq(1,365,by=5)
x1 \leftarrow c(1,3,4)
x2 < -c(2,3,4)
# Function that returns kernel class object
Matern32 <- function(sigmaf = 1, ell = 1)</pre>
 rval <- function(x1,x2){</pre>
   n1 \leftarrow length(x1)
   n2 \leftarrow length(x2)
   K <- matrix(NA,n1,n2)</pre>
    for (i in 1:n2){
      K[,i] \leftarrow sigmaf^2*exp(-0.5*((x1-x2[i])/ell)^2)
    }
    return(K)
  class(rval) <- "kernel"</pre>
 return(rval)
}
MaternFunc = Matern32(sigmaf = 1, ell = 2)
```

```
#evaluate it in the point x = 1; x' = 2
cat("The kernel between points (1,2) is :", MaternFunc(1,2))
cat("\n")
cat("\n")
# use the kernelMatrix function to compute the covariance matrix K(X;X*)
covM <- kernelMatrix(MaternFunc, x1,x2)</pre>
cat("The kernel between points (1,3,4) and (2,3,4) is:")
cat("\n")
covM
# Subset data
temp <- dataQ2$temp[Stime]</pre>
# Find the noise value
polyFit <- lm(temp ~ Stime + Stime^2)</pre>
sigmaNoise = sd(polyFit$residuals)
MaternFunc2 = Matern32(sigmaf = 20, ell = 0.2)
# Fit the GP with built in Square expontial kernel (called rbfdot in kernlab)
GPfit <- gausspr(Stime, temp, kernel = MaternFunc2, var = sigmaNoise^2)</pre>
postMean2 <- predict(GPfit,Stime)</pre>
plot(Stime, temp, ylim=c(-30,40),xlab="time",ylab="temperature")
lines(Stime,postMean2, col="red", lwd = 2)
legend("topright",c("Observations","Post. Mean"),
        col = c("black", "red"), pch = c(1,-1), lty = c(0, 1), lwd=c(0,2))
x<-scale(Stime)
xs<-scale(Stime) # XStar
n <- length(x)
SEkernel <- MaternFunc2
Kss <- kernelMatrix(kernel = SEkernel, x = xs, y = xs)</pre>
Kxx <- kernelMatrix(kernel = SEkernel, x = x, y = x)</pre>
Kxs <- kernelMatrix(kernel = SEkernel, x = x, y = xs)</pre>
Covf = Kss-t(Kxs)%*%solve(Kxx + sigmaNoise^2*diag(n), Kxs) # Covariance matrix of fStar
plot(Stime, temp, ylim=c(-30,65),xlab="time",ylab="temperature")
lines(Stime,postMean2, col="red", lwd = 2)
# Probability intervals for fStar
lines(Stime, postMean2 - 1.96*sqrt(diag(Covf)), col = "green")
lines(Stime, postMean2 + 1.96*sqrt(diag(Covf)), col = "green")
# Prediction intervals for yStar
lines(Stime, postMean2 - 1.96*sqrt((diag(Covf) + sigmaNoise^2)), col = "purple")
lines(Stime, postMean2 + 1.96*sqrt((diag(Covf) + sigmaNoise^2)), col = "purple")
legend("topright",c("Observations","Post. Mean","Prob. bands","Pred. bands"),
        col = c("black", "red", "green", "purple"), pch = c(1,-1,-1,-1), lty = c(0, 1,1,1), lwd=c(0,2,2,2)
# Find the noise value
polyFit4 <- lm(temp ~ rep.int(Sday,6) )</pre>
sigmaNoise4 = sd(polyFit4$residuals)
# Fit the GP with built in Square expontial kernel (called rbfdot in kernlab)
GPfit4 <- gausspr(rep.int(Sday,6), temp, kernel = MaternFunc2, var = sigmaNoise4^2)</pre>
postMean4 <- predict(GPfit4,rep.int(Sday,6))</pre>
```

```
plot(Stime, temp,xlab="time",ylab="temperature",ylim=c(-30,45))
lines(Stime,postMean2, col="red", lwd = 2)
lines(Stime,postMean4, col="blue", lwd = 2)
legend("topright",c("Observations","Post. Mean(time)","Post. Mean(day)"),
        col = c("black","red", "blue"), pch = c(1,-1,-1), lty = c(0, 1,1), lwd=c(0,2,2))
# Function that returns kernel class object
Matern5 <- function(sigmaf, ell1, ell2, d){</pre>
  val <- function(x1, x2){</pre>
    n1 \leftarrow length(x1)
    n2 \leftarrow length(x2)
    K <- matrix(NA,n1,n2)</pre>
    for (i in 1:n2){
      K[,i] < - sigmaf^2*exp(-2*sin(pi*abs(x1-x2[i])/d)^2/ell1^2)*exp(-0.5*abs(x1-x2[i])^2/ell2^2)
    }
    return(K)
  class(val) <- "kernel"</pre>
 return(val)
MaternFunc5 <- Matern5(20, 1, 10, 365/sd(time))
GPfit5 <- gausspr(Stime, temp, kernel = MaternFunc5, var = sigmaNoise^2)</pre>
postMean5 <- predict(GPfit5,Stime)</pre>
plot(Stime, temp,ylim=c(-30,45),xlab="time",ylab="temperature")
lines(Stime,postMean2, col="red", lwd = 2)
lines(Stime,postMean4, col="blue", lwd = 2)
lines(Stime,postMean5, col="green", lwd = 2)
legend("topright",c("Observations","Post. Mean(time)","Post. Mean(day)","Post. Mean(periodic)"),
        col = c("black","red","blue","green"), pch = c(1,-1,-1,-1), lty = c(0, 1,1,1), lwd=c(0,2,2,2))
# read data from github
data <- read.csv("https://github.com/STIMALiU/AdvMLCourse/raw/master/GaussianProcess/Code/banknoteFraud
                  header=FALSE, sep=",")
names(data) <- c("varWave", "skewWave", "kurtWave", "entropyWave", "fraud")</pre>
data[,5] <- as.factor(data[,5])</pre>
head(data)
set.seed(111);
SelectTraining <- sample(1:dim(data)[1], size = 1000,
                                          replace = FALSE)
training <- data[SelectTraining,]</pre>
testing <- data[-SelectTraining,]</pre>
accuracy <- function (conf_mat){(conf_mat[1,1] + conf_mat[2,2]) / sum(conf_mat)}</pre>
GPfitFraud <- gausspr(fraud ~ varWave + skewWave, data=training)</pre>
# class probabilities
probPreds <- predict(GPfitFraud, training, type="probabilities")</pre>
x1 <- seq(min(training[,1]),max(training[,1]),length=100)</pre>
x2 <- seq(min(training[,2]),max(training[,2]),length=100)
gridPoints <- meshgrid(x1, x2)</pre>
gridPoints <- cbind(c(gridPoints$x), c(gridPoints$y))</pre>
gridPoints <- data.frame(gridPoints)</pre>
```

```
names(gridPoints) <- names(training)[1:2]</pre>
probPreds <- predict(GPfitFraud, gridPoints, type="probabilities")</pre>
# predict on the testing set
pred1 <- predict(GPfitFraud, training)</pre>
conM1 <- table("Prediction"=pred1, "True"=training$fraud) # confusion matrix</pre>
conM1
# Plotting for Prob(setosa)
contour(x1,x2,matrix(probPreds[,1],100,byrow = TRUE), 20, xlab = "varWave",
        ylab = "skewWave", main = 'Contour of Prediction Probabilities')
points(training[training[,5]==0,1],training[training[,5]==0,2],col="blue")
points(training[training[,5]==1,1],training[training[,5]==1,2],col="red")
legend("bottomright", legend=c("0","1"), pch=c(20,20), col=c("red","blue"))
# predict on the testing set
predT <- predict(GPfitFraud, testing)</pre>
conM2 <- table("Prediction"=predT, "True"=testing$fraud) # confusion matrix</pre>
conM2
GPfitFraud3 <- gausspr(fraud ~ ., data=training)</pre>
# predict on the testing set
pred3 <- predict(GPfitFraud3,testing)</pre>
conM3 <- table("Prediction"=pred3, "True"= testing$fraud) # confusion matrix</pre>
conM3
```