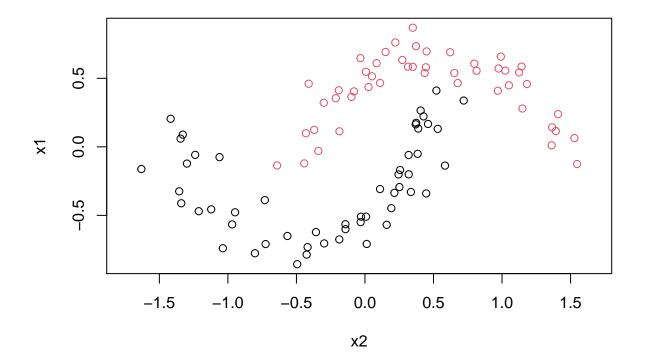
ST340 Lab 7: Support vector machines

2020-21

Load package e1071:

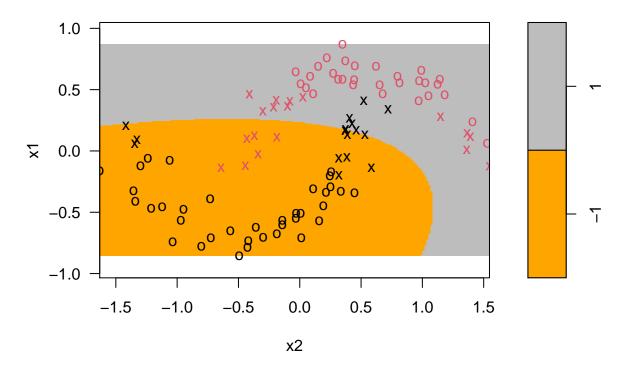
(a) Run the following code, and then construct and plot an SVM with a quadratic kernel. What is its test set accuracy? (The code to answer this is on the lecture slides).

```
set.seed(1)
th=runif(100,-pi,pi)
y=sign(th)
x1=sin(th)-y/3+rnorm(100,sd=0.1)
x2=cos(th)+y/2+rnorm(100,sd=0.1)
plot(x2,x1,asp=1,col=(y+3)/2)
```



```
d <- data.frame(x1=x1,x2=x2,y=factor(y))
s = svm(y ~., d, kernel="polynomial", degree=2, coef0 = 1, cost = 1)  # Default cost is 1
paste("accuracy", mean(predict(s,d[,1:2])==y),"  ||  ", "Number of support vectors:", nrow(s$SV))
## [1] "accuracy 0.87  || Number of support vectors: 34"</pre>
```

SVM classification plot

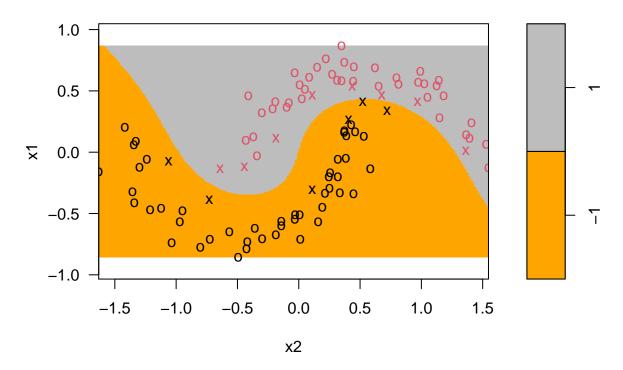


(b) Use leave-one-out-cross-validation to compare polynomial kernels. Write some code to perform a grid search over possible choices of parameters. For simplicity, fix c=1 (coef0 = 1) and C=1 (cost=1) and search over the remaining parameters γ and p and to select the model which is optimal in terms of cross-validation error. A suggested grid is:

```
p.range <- 1:10
gamma.range <- c(0.001,0.01,0.1,0.5,1,2,5,10)
CV_grid = matrix(NA,length(p.range),length(gamma.range))
rownames(CV_grid) = paste("P =",p.range)
colnames(CV_grid) = paste("Gamma =",gamma.range)

for(i in 1:length(p.range)){
    for(j in 1:length(gamma.range)){
        s = svm(y ~., d, kernel="polynomial", degree = p.range[i], gamma = gamma.range[j], coef0 = 1, cross = CV_grid[i,j] = s$tot.accuracy
    }
}
# From this LOOCV it seems that setting Gamma = 1 and P = 3 performs the best
s = svm(y ~., d, kernel="polynomial", degree = 3, gamma = 1, coef0 = 1)
plot(s,d, asp=1, grid = 200,col = c("orange","grey74"))</pre>
```

SVM classification plot



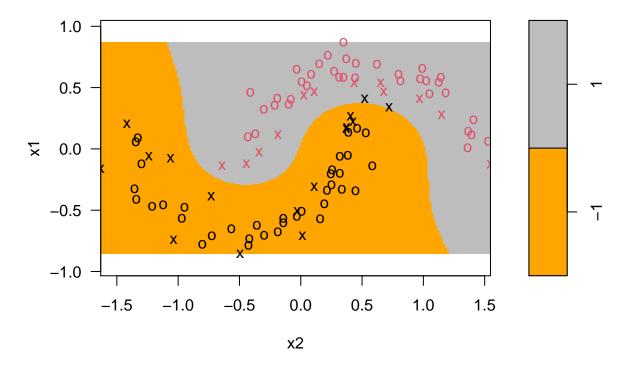
(Hint: You do not need to write a function to compute the cross-validation error: look at the cross flag in ?svm. You can extract the accuracies of the cross-validation sets from an svm object s using s\$accuracies and s\$tot.accuracy.)

(c) Repeat part (b) to choose a value of γ for the radial basis function.

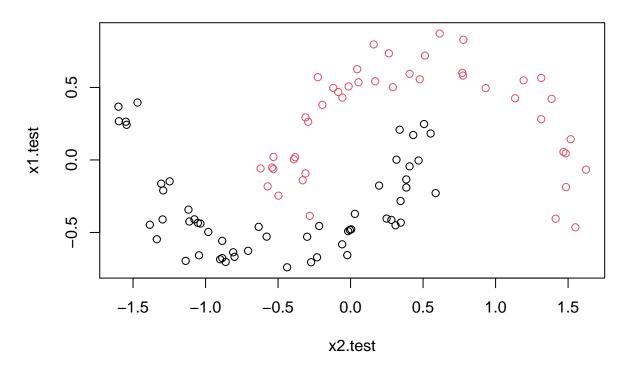
```
gamma.range \leftarrow c(0.001,0.01,0.1,0.5,1,2,5,10)
CV_vector = matrix(NA, nrow = 1, ncol = length(gamma.range))
colnames(CV_vector) = gamma.range
for(j in 1:length(gamma.range)){
  s = svm(y ~., d, kernel="radial", gamma = gamma.range[j], cost = 1, cross = nrow(d))
  CV_vector[1,j] = s$tot.accuracy
}
print(CV_vector)
##
        0.001 0.01 0.1 0.5 1 2 5 10
           52
                83 85 97 98 98 98 98
## [1,]
colnames(CV_vector)[which.max(CV_vector)]
## [1] "1"
# From this LOOCV it seems that setting Gamma = 1 performs the best
# After further investigation, doesn't seem that we can get better prediction accuracy than 98%
```

```
s = svm(y ~., d, kernel="radial", gamma = 1)
plot(s,d, asp=1, grid = 200,col = c("orange", "grey74"))
```

SVM classification plot



(d) Compare the performance of your chosen polynomial kernel and RBF kernel by simulating an independent test dataset and evaluating their accuracies:



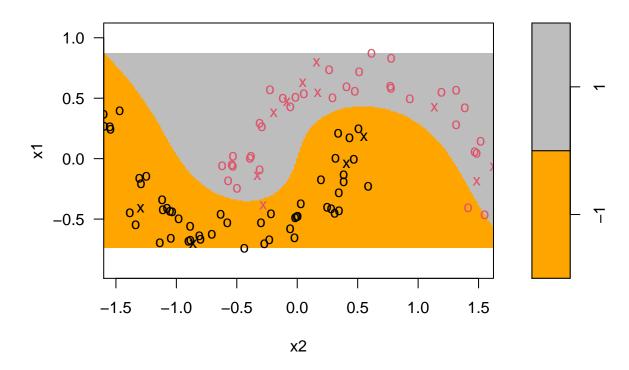
```
d.test <- data.frame(x1=x1.test,x2=x2.test,y=factor(y.test))
mean(predict(s.poly,d.test)==y.test)

## [1] 0.98
mean(predict(s.rbf,d.test)==y.test)</pre>
```

[1] 0.99

plot(s.poly,d.test,grid=200,asp=1,col = c("orange","grey74"), main = "polynomial Kernel\nDegree = 3, Gar

SVM classification plot



plot(s.rbf,d.test,grid=200,asp=1, col = c("orange","grey74"))

SVM classification plot

