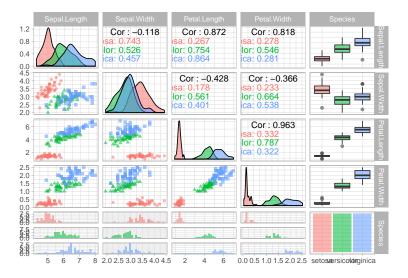
1 Classifier

1.1 proof

$$\begin{split} g(x) &= \langle C_+ - C_-, X - C \rangle = \langle C_+, X \rangle - \langle C_-, X \rangle - \langle C_+, C \rangle + \langle C_-, C \rangle \\ \langle C_+, X \rangle &= \frac{1}{n_+} \sum_{i \in I_+}^n \langle x_i, x \rangle; \\ \langle C_-, X \rangle &= \frac{1}{n_-} \sum_{i \in I_-}^n \langle x_i, x \rangle; \\ \langle C_+, C \rangle &= \langle C_+, \frac{1}{2}C_+ \rangle + \langle C_+, \frac{1}{2}C_- \rangle = \frac{1}{2n_+^2} \sum_{(i,j) \in I_+} \langle x_i, x_j \rangle + \frac{1}{2} \langle C_+, C_- \rangle \\ \langle C_-, C \rangle &= \langle C_-, \frac{1}{2}C_+ \rangle + \langle C_-, \frac{1}{2}C_- \rangle = \frac{1}{2} \langle C_+, C_- \rangle + \frac{1}{2n_-^2} \sum_{(i,j) \in I_-} \langle x_i, x_j \rangle \\ g(x) &= \frac{1}{n_+} \sum_{i \in I_+}^n \langle x_i, x \rangle - \frac{1}{n_-} \sum_{i \in I_-}^n \langle x_i, x \rangle - \frac{1}{2n_+^2} \sum_{(i,j) \in I_+} \langle x_i, x_j \rangle - \frac{1}{2} \langle C_+, C_- \rangle + \frac{1}{2} \langle C_+, C_- \rangle + \frac{1}{2n_-^2} \sum_{(i,j) \in I_-} \langle x_i, x_j \rangle \\ &= \sum_{i=1}^n \alpha_i \langle x_i, x \rangle + b \end{split}$$
 where $\alpha_i = \begin{cases} \frac{1}{n_+} & y_i = +1 \\ -\frac{1}{n_-} & y_i = -1 \end{cases}; \ b = \frac{1}{2n_-^2} \sum_{(i,j) \in I_-} \langle x_i, x_j \rangle - \frac{1}{2n_+^2} \sum_{(i,j) \in I_+} \langle x_i, x_j \rangle$

1.2 iris data clasification

• Import the iris data

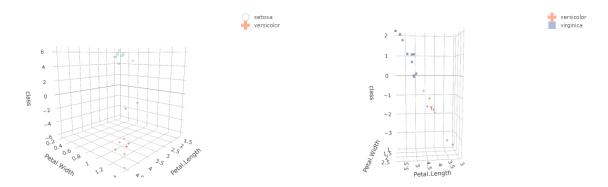


```
# Define the train and test sets
iris$class <- NA
iris_setosa <- iris[iris$Species=="setosa",]
iris_versicolor <- iris[iris$Species=="versicolor",]
iris_virginica <- iris[iris$Species=="virginica",]
iris_train_se<- iris_setosa[1:40,]
iris_train_ve<- iris_versicolor[1:40,]
iris_train_vi<- iris_virginica[1:40,]
iris_test_se<- iris_setosa[41:50,]
iris_test_ve<- iris_versicolor[41:50,]
iris_test_ve<- iris_versicolor[41:50,]
iris_train_se.ve<- rbind(iris_train_se,iris_train_ve)
iris_train_ve.vi<- rbind(iris_train_ve,iris_train_vi)
iris_test_se.ve<- rbind(iris_test_se,iris_test_ve)
iris_test_ve.vi<- rbind(iris_test_se,iris_test_ve)
iris_test_ve.vi<- rbind(iris_test_se,iris_test_ve)</pre>
```

```
# Define the kernel function and Computing the classifier
k = function(x,y) return(sum(x*y))
# setosa v.s.versicolor
k.pp=outer(1:40,1:40,Vectorize(function(i,j) k(iris_train_se[i,1:4],iris_train_se[j,1:4])))
k.mm=outer(1:40,1:40,Vectorize(function(i,j) k(iris_train_ve[i,1:4],iris_train_ve[j,1:4])))
b=(sum(k.mm)/(40^2)-sum(k.pp)/(40^2))/2
alpha=ifelse(iris_train_se.ve$Species=="setosa",1/40,-1/40)
k.x=outer(1:80,1:20,Vectorize(function(i,j) k(iris_train_se.ve[i,1:4],iris_test_se.ve[j,1:4])))
iris_test_se.ve[,6]=(t(k.x)%*%alpha+b)
```

```
# virginica v.s.versicolor
k.pp=outer(1:40,1:40,Vectorize(function(i,j) k(iris_train_vi[i,1:4],iris_train_vi[j,1:4])))
k.mm=outer(1:40,1:40,Vectorize(function(i,j) k(iris_train_ve[i,1:4],iris_train_ve[j,1:4])))
b=(sum(k.mm)/(40^2)-sum(k.pp)/(40^2))/2
alpha=ifelse(iris_train_ve.vi$Species="virginica",1/40,-1/40)
k.x=outer(1:80,1:20,Vectorize(function(i,j) k(iris_train_ve.vi[i,1:4],iris_test_ve.vi[j,1:4])))
iris_test_ve.vi[,6]=(t(k.x)%*%alpha+b))
# Evaluate the classifier
iris_test_se.ve$evaluate=ifelse(iris_test_se.ve$class>0,"setosa","versicolor")
iris_test_se.ve$evaluate=ifelse(iris_test_se.ve$Species==iris_test_se.ve$evaluate,"Rigt","Wrong")
error_rate1 <- length(which(iris_test_se.ve$evaluate=="Wrong"))/20
iris_test_ve.vi$evaluate=ifelse(iris_test_ve.vi$Cpecies==iris_test_ve.vi$evaluate,"Rigt","Wrong")
error_rate2 <- length(which(iris_test_ve.vi$evaluate=="Wrong"))/20</pre>
```

Error rate = 0% and 5% in two tests respectively.



The left figure of setosa v.s. versicolor shows that the two species are separated by zero plane, while virginica v.s. versicolor shows that a few points are close to zero. The tables also show a misclassified point in virginica v.s. versicolor.

Table 1: Confusion matrix

	Actural Species								
	test 1	Setosa	Versicolor	test 2	Virginica	Versicolor			
Test Species	Setosa	10	0	Virginica	9	0			
	Versicolor	0	10	Versicolor	1	10			

Table 2: setosa v.s.versicolor//virginica v.s.versicolor

Species	class	evaluate	Species1	class1	evaluate1
setosa	5.856	Rigt	versicolor	-1.575	Rigt
setosa	5.536	Rigt	versicolor	-0.7495	Rigt
setosa	6.35	Rigt	versicolor	-1.907	Rigt
setosa	4.665	Rigt	versicolor	-3.482	Rigt
setosa	4.135	Rigt	versicolor	-1.69	Rigt
setosa	5.429	Rigt	versicolor	-1.638	Rigt
setosa	5.215	Rigt	versicolor	-1.592	Rigt
setosa	5.869	Rigt	versicolor	-1.157	Rigt
setosa	5.239	Rigt	versicolor	-3.708	Rigt
setosa	5.548	Rigt	versicolor	-1.739	Rigt
versicolor	-5.097	Rigt	virginica	1.566	Rigt
versicolor	-6.206	Rigt	virginica	0.9795	Rigt
versicolor	-4.246	Rigt	virginica	-0.02223	Wrong
versicolor	-1.446	Rigt	virginica	1.968	Rigt
versicolor	-4.667	Rigt	virginica	1.795	Rigt
versicolor	-4.451	Rigt	virginica	0.968	Rigt
versicolor	-4.63	Rigt	virginica	0.119	Rigt
versicolor	-5.402	Rigt	virginica	0.6535	Rigt
versicolor	-0.6631	Rigt	virginica	0.9918	Rigt
versicolor	-4.411	Rigt	virginica	0.02902	Rigt

2 Perceptron

2.1 pseduo code

```
Init:

- define the classifier f(x) = w^T x

- define perceptron algorithm

- set the sample size =n

- set initial weight with w \leftarrow y_1 x_1

- set initial misclassified argument with TRUE

While misclassified=TURE

- change argument with FALSE

- For i = 2 \dots n do

— If y_i w^T x_i < 0 then w \leftarrow w + y_i x_i

— change argument with TRUE

— EndIf

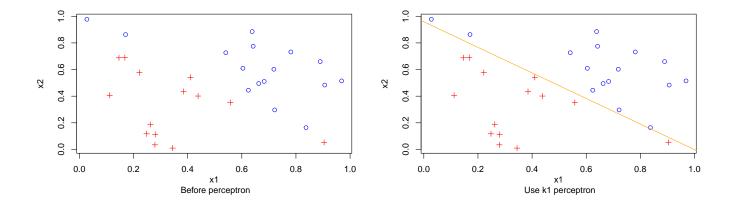
- EndFor

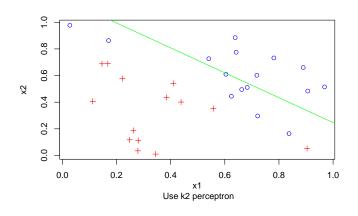
EndWhile
```

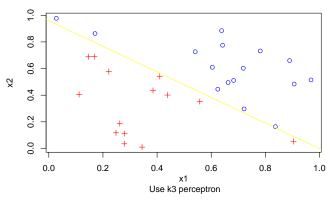
2.2 Using 3 Kernels

```
30; dim <- 2; threshold <-1
                                                                   ## generate 2d data
   <- runif(n * dim)
<- matrix(x, ncol = dim)
<- ifelse(apply(x, 1, sum) < threshold, -1, 1)</pre>
data <- cbind(x, y, alpha = rep(1, n))
k<- function(x,w){return(sum(x*w))}
k_perceptron <- function(data,dim) {</pre>
                                                                   ## define the kernel function
                                                                   ## difine perceptron algorithm
   <- nrow(data)
                                                                   ## get the sample size
## select y
## initialize weight with y_1*x_1
                                                                   ## initialize the argument
                                                                   ## start to loop
                                                                   ## loop times = sample size-1
         if (y[i]*k(data[i,-dim-1],w) < 0) { ## if expectation does not match y w <- w + y[i] * data[i,-dim-1] ## update the weights misclassfied <- TRUE ## reset argument and check next one
fk <- function(x,y) {return(sum(x*y))} ## Use k1
wut <- k_perceptron(data,2))</pre>
## 4.004353 4.173336 -4.000000
k <- function(x,y) {return(sum(x*y)+1)} ## Use k2
(w2 <- k_perceptron(data,2))
## 3.957389 4.230957 -5.000000
k <- function(x,y) {return(sign(x,*,y))} ## Use k3
(w3 <- k_perceptron(data,2))</pre>
    4.004353 4.173336 -4.000000
```

The output shows that the perceptrons with $k = x^T y$ and $k = \operatorname{sign}(x^T y)$ can classify the data. The $k = x^T y + 1$ is biased.







3 Kernels over $\mathcal{X} = \mathbb{R}^2$

 $x = (x_1, x_2) \in \mathbb{R}^2 \text{ and } y = (y_1, y_2) \in \mathbb{R}^2,$

3.1 Let $\phi(x) = (x_1^2, \sqrt{2}x_1x_2, x_2^2)$, verify that $\phi(x)^T \phi(y) = (x^T y)^2$

$$(x^T y)^2 = (\begin{bmatrix} x_1 \\ x_2 \end{bmatrix}_{2 \times 1} \begin{bmatrix} y_1 & y_2 \end{bmatrix}_{1 \times 2})^2 = (x_1 y_1 + x_2 y_2)^2 = x_1^2 y_1^2 + 2x_1 x_2 y_1 y_2 + x_2^2 y_2^2$$

$$= \begin{bmatrix} x_1^2 \\ \sqrt{2} x_1 x_2 \\ x_2^2 \end{bmatrix}_{3 \times 1} \begin{bmatrix} y_1^2 & \sqrt{2} y_1 y_2 & y_2^2 \end{bmatrix}_{1 \times 3} = \phi(x)^T \phi(y) \qquad \blacksquare$$

3.2 Find a function $\phi(x): \mathbb{R}^2 \mapsto \mathbb{R}^6$ such that for any $(x,y), \ \phi(x)^T \phi(y) = (x^Ty+1)^2$

$$(x^{T}y+1)^{2} = (x_{1}y_{1} + x_{2}y_{2} + 1)^{2} = x_{1}^{2}y_{1}^{2} + 2x_{1}y_{1} + 2x_{1}x_{2}y_{1}y_{2} + 2x_{2}y_{2} + x_{2}^{2}y_{2}^{2} + 1$$

$$= \begin{bmatrix} x_{1}^{2} \\ \sqrt{2}x_{1} \\ \sqrt{2}x_{1}x_{2} \\ \sqrt{2}x_{2} \\ x_{2}^{2} \\ 1 \end{bmatrix}_{6\times 1} [y_{1}^{2} \sqrt{2}y_{1} \sqrt{2}y_{1}y_{2} \sqrt{2}y_{2} y_{2}^{2} 1]_{1\times 6}$$

Thus,

$$\phi(x) = (x_1^2, \sqrt{2}x_1, \sqrt{2}x_1x_2, \sqrt{2}x_2, x_2^2, 1)$$

3.3 Find a function $\phi(x): \mathbb{R}^2 \to \mathbb{R}^9$ such that for any (x,y), $\phi(x)^T \phi(y) = (x^T y + 1)^2$

$$(x^{T}y+1)^{2} = x_{1}^{2}y_{1}^{2} + 2x_{1}y_{1} + \frac{1}{2}x_{1}x_{2}y_{1}y_{2} + \frac{1}{2}x_{2}x_{1}y_{2}y_{1} + \frac{1}{2}x_{1}^{0.5}x_{2}x_{1}^{0.5}y_{1}^{0.5}y_{2}y_{1}^{0.5} + \frac{1}{2}x_{2}^{0.5}x_{1}x_{2}^{0.5}y_{2}^{0.5}y_{1}y_{2}^{0.5} + 2x_{2}y_{2} + x_{2}^{2}y_{2}^{2} + 1$$

$$=\begin{bmatrix} x_1^2 \\ \sqrt{2}x_1 \\ \frac{1}{\sqrt{2}}x_1x_2 \\ \frac{1}{\sqrt{2}}x_2x_1 \\ \frac{1}{\sqrt{2}}x_1^{0.5}x_2x_1^{0.5} \\ \frac{1}{\sqrt{2}}x_2^{0.5}x_1x_2^{0.5} \\ \sqrt{2}x_2 \\ x_2^2 \\ 1 \end{bmatrix}_{9\times 1} \begin{bmatrix} y_1^2 & \sqrt{2}y_1 & \frac{1}{\sqrt{2}}y_1y_2 & \frac{1}{\sqrt{2}}y_2y_1 & \frac{1}{\sqrt{2}}y_1^{0.5}y_2y_1^{0.5} & \frac{1}{\sqrt{2}}y_2^{0.5}y_1y_2^{0.5} & \sqrt{2}y_2 & y_2^2 & 1 \end{bmatrix}_{1\times 9}$$

Thus,

$$\phi(x) = (x_1^2, \sqrt{2}x_1, \frac{1}{\sqrt{2}}x_1x_2, \frac{1}{\sqrt{2}}x_2x_1, \frac{1}{\sqrt{2}}x_1^{0.5}x_2x_1^{0.5}, \frac{1}{\sqrt{2}}x_2^{0.5}x_1x_2^{0.5}, \sqrt{2}x_2, x_2^2, 1)$$

Verify that $K(x,y) = (1+x^Ty)^d$ for d=1,2... is a positive definite kernel

We know that
$$x^T y = \begin{bmatrix} x_1 \\ x_2 \end{bmatrix}_{2 \times 1} [y_1 \quad y_2]_{1 \times 2} = x_1 y_1 + x_2 y_2 = y^T x$$
, then

$$k(x,y) = (1 + x^T y)^d = (1 + y^T x)^d = k(y,x)$$

Thus, k(x, y) is symmetric.

$$\exists \phi(x) = \left(\sqrt{\binom{k}{0}} x_1^k x_2^0, \sqrt{\binom{k}{1}} x_1^{k-1} x_2^1, \cdots, \sqrt{\binom{k}{k-1}} x_1^1 x_2^{k-1}, \sqrt{\binom{k}{k}} x_1^0 x_2^k\right)$$

Make

$$k(\phi(x), \phi(y)) = \phi(x)^T \phi(y) = \sum_{l=0}^k \binom{k}{l} (x_1 y_1)^{k-l} (x_2 y_2)^l = (x_1 y_1 + x_2 y_2)^k = (x^T y)^k$$

 $k(\phi(x), \phi(y))$ is a p.d. kernel and $\|\sum_{i=1}^{2} \alpha_i \phi_i(x)\|^2 \ge 0$. Therefore,

$$k(x,y) = (1 + x^T y)^d = \sum_{k=0}^d \binom{d}{k} (x^T y)^k 1^{d-k} = \sum_{k=0}^d \binom{d}{k} \phi(x)^T \phi(y)$$

For $d, k \in \mathbb{N}^+$,

$$\sum_{i=1}^{2} \sum_{j=1}^{2} \alpha_{i} \alpha_{j} \langle x_{i}, x_{j} \rangle_{R^{d}} = \sum_{i=1}^{2} \sum_{j=1}^{2} \alpha_{i} \alpha_{j} \left[\sum_{k=0}^{d} \binom{d}{k} \phi_{i}(x)^{T} \phi_{j}(x) \right] = \sum_{k=0}^{d} \binom{d}{k} \left[\sum_{i=1}^{2} \sum_{j=1}^{2} \alpha_{i} \alpha_{j} \langle \phi_{i}(x)^{T} \phi_{j}(x) \rangle_{R} \right]$$

$$= \sum_{k=0}^{d} \binom{d}{k} \| \sum_{i=1}^{2} \alpha_{i} \phi_{i}(x) \|^{2} \ge 0$$

Therefore, for $x_1, x_2, x_i \in \mathbb{R}$; $\alpha_1, ..., \alpha_2, \alpha_i \in \mathbb{R}$ $k(x, y) = (1 + x^T y)^d$ is a positive definite kernel.

find a function $\phi:\mathbb{R}^2 \mapsto H$, where H is an inner product space such that for any (x,y), $\langle \phi(x), \phi(y) \rangle_H = x^T y - 1$

$$k(x,y) = \langle \phi(x), \phi(y) \rangle_H = x^T y - 1 = \begin{bmatrix} x_1 \\ x_2 \end{bmatrix}_{2 \times 1} [y_1 \quad y_2]_{1 \times 2} - 1 = x_1 y_1 + x_2 y_2 - 1$$

For all $|x| < \frac{\sqrt{2}}{2}$, when $\alpha_1 = \alpha_2 = 1$, $\langle \phi(x), \phi(x) \rangle_H = 2x^2 - 1 < 0$. By the property of Kernel, k(x,y) should be positive.

Therefore, there isn't a function $\phi: \mathbb{R}^2 \to H$ that can make $\langle \phi(x), \phi(y) \rangle_H = x^T y - 1$.