

13강

분류분석 (1)

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1 판별분석



붓꽃

setosa



versicolor



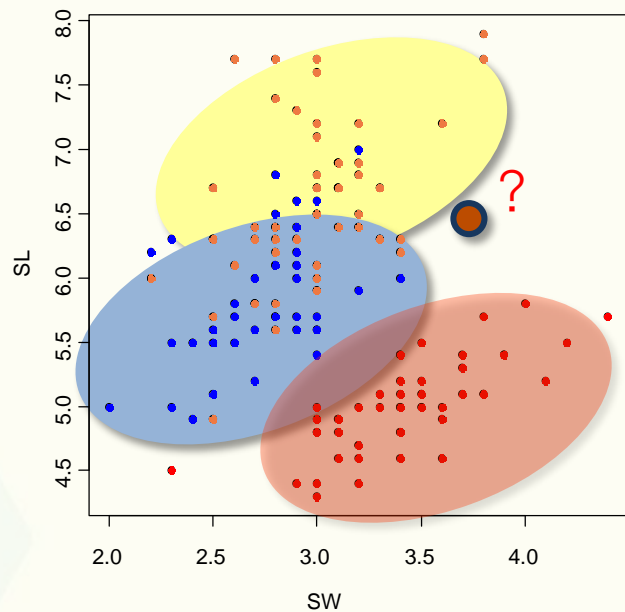
virginica



?



판별분석



판별분석

판별분석 (Discriminant Analysis) :

예 : $D = 0.83*SL + 1.53*SW - 2.2*PL - 2.8*PW$

만약 $D > 0$ 이면 "세토사"

선형판별분석(Linear DA) : 판별식 D가 독립변수들에 대하여 선형

이차판별분석(Quadratic DA) : 판별식 D가 독립변수들에 대하여 2차식



판별분석의 기본 틀

종속변수 : $y \in \{y_{(0)}, \dots, y_{(K)}\}$ 예 : 붓꽃의 종류

독립변수 : $\mathbf{x} = (x_1, x_2, \dots, x_p)'$ 예 : (꽃잎, 꽃받침) × (폭, 길이)

$$\mathbf{x}|y_{(k)} \sim f(\mathbf{x}|y_{(k)}) \quad k = 0, 1, \dots, K$$

$$\hat{k} = \arg \max_k f(y_{(k)}|\mathbf{x}) = \arg \max_k \log f(y_{(k)}|\mathbf{x})$$

$$\log f(y_{(k)}|\mathbf{x}) \propto \log f(\mathbf{x}|y_{(k)}) + \log f(y_{(k)}) \quad \text{베이즈 정리}$$

$$\hat{y} = y_{(\hat{k})}$$



베이즈 정리

$$x|y \sim f(x|y) \quad \longrightarrow \quad y|x \sim f(y|x)$$

$$f(x, y) = f(x) f(y|x) = f(x|y) f(y)$$

$$f(x) = \sum_y f(x|y) f(y)$$

$$f(y|x) = \frac{f(x|y) f(y)}{f(x)}$$

$$f(y|x) \propto f(x|y) f(y) \quad \log f(y|x) \propto \log f(x|y) + \log f(y)$$



정규분포 가정

$$\mathbf{x}|y_{(k)} \sim N_p(\mu_k, \Sigma_k) \quad k = 0, 1, \dots, K$$

$$\mu_k = (\mu_{k1}, \mu_{k2}, \dots, \mu_{kp})' \quad \Sigma_k = \begin{pmatrix} \sigma_{k11} & \dots & \sigma_{k1p} \\ \dots & \dots & \dots \\ \sigma_{kp1} & \dots & \sigma_{kpp} \end{pmatrix}$$


$$Q_k(\mathbf{x}) = -2 \log f(y_{(k)}|\mathbf{x})$$

$$= -2 \log f(\mathbf{x}|y_{(k)}) - 2 \log f(y_{(k)}) + \text{constant}$$


$$= (\mathbf{x} - \mu_k)' \Sigma_k^{-1} (\mathbf{x} - \mu_k) + \log |\Sigma_k| - 2 \log f(y_{(k)}) \quad \downarrow$$

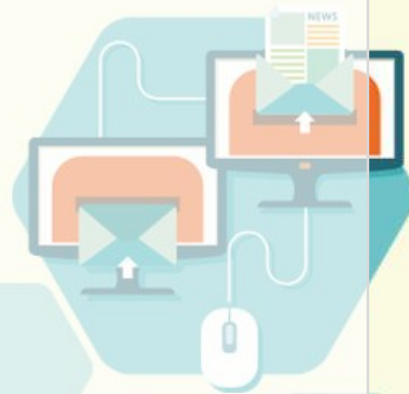


LDA & QDA : 정규분포

QDA : $Q_k(\mathbf{x}) = (\mathbf{x} - \mu_k)' \Sigma_k^{-1} (\mathbf{x} - \mu_k) + \log |\Sigma_k| - 2 \log f(y_{(k)})$ 

LDA : $\Sigma_k = \Sigma \quad k = 0, 1, \dots, K$

$$L_k = \mathbf{x}' \Sigma^{-1} \mu_k - (1/2) \mu_k' \Sigma^{-1} \mu_k + \log f(y_{(k)})$$
 



LDA : 2 그룹

LDA : $\Sigma_k = \Sigma \quad k = 0, 1, \dots, K$

$$L_k = \mathbf{x}'\Sigma^{-1}\mu_k - (1/2)\mu_k'\Sigma^{-1}\mu_k + \log f(y_{(k)}) \quad \uparrow$$

두 그룹의 경우 ($K = 1$) 선형판별식 :

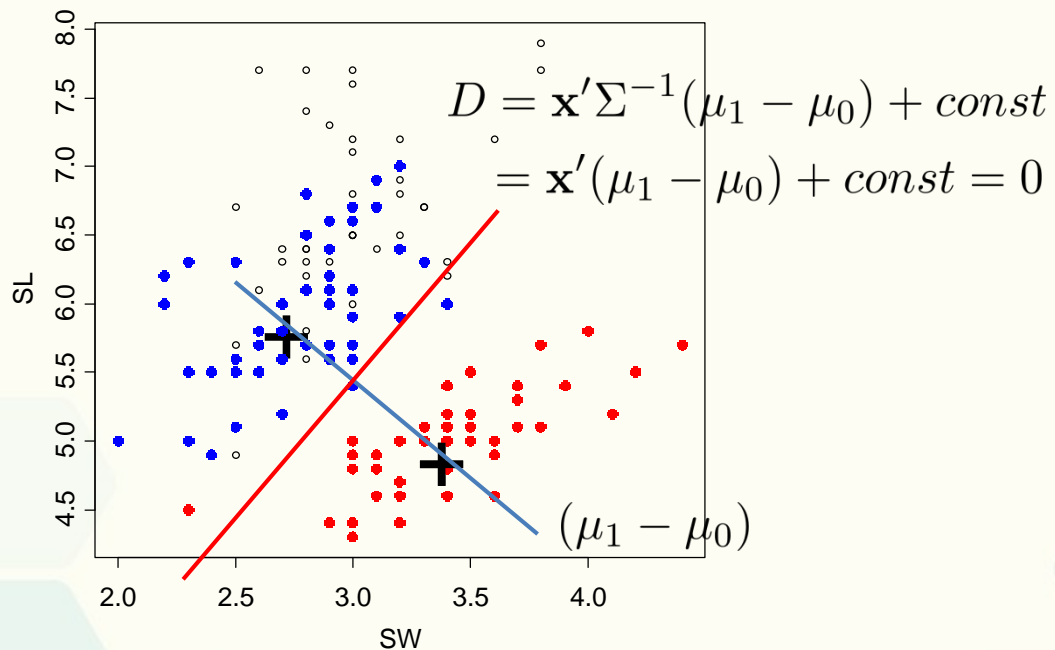
$$D = L_1 - L_0 = \mathbf{x}'\Sigma^{-1}(\mu_1 - \mu_0) + constant$$

- $D > c_0 (= 0) \Rightarrow \hat{k} = 1$
- $D < c_0 (= 0) \Rightarrow \hat{k} = 0$



LDA : 2 그룹

단순화하여 $\Sigma = I$ 라고 가정하고,
setosa 와 versicolor 경우.



LDA : 다그룹

그룹내 분산 : $\Sigma = E [Var(\mathbf{x}|y)]$

그룹간 분산 : $\Sigma_b = Var [E(\mathbf{x}|y)]$

$D = w' \mathbf{x}$ 라 하면,

$$S = \frac{Var [E(w' \mathbf{x}|y)]}{E [Var(w' \mathbf{x}|y)]} = \frac{w' \Sigma_b w}{w' \Sigma w} \quad \uparrow$$

$$D_k = w'_k \mathbf{x} \quad k = 1, \dots, K$$

w_k : $\Sigma^{-1} \Sigma_b$ 의 고유벡터들(eigen vectors) .



LDA와 LRA

$$p_k = f(y_{(k)}|\mathbf{x}) \quad k = 0, 1, \dots, K$$

LRA : 로지스틱 회귀분석(Logistic Regression Analysis)

$$(y_{(0)}, y_{(1)}, \dots, y_{(K)}) \sim Multinom(n, p_0, \dots, p_K)$$

$$\log(p_k/p_0) = \mathbf{x}'\beta_k + \alpha_k$$

LDA : $\mathbf{x}|y_{(k)} \sim N_p(\mu_k, \Sigma)$ **x : 양적변수**

$$\log(p_k/p_0) = \log f(\mathbf{x}|y_{(k)}) - \log f(\mathbf{x}|y_{(0)}) + const$$

$$= \mathbf{x}'\Sigma^{-1}(\mu_1 - \mu_0) + const$$

$$= \mathbf{x}'\beta_k + \alpha_k \quad \beta_k = (\beta_{k1}, \dots, \beta_{kp})'$$



2 판별분석의 적용 예



R의 lda 와 qda 함수

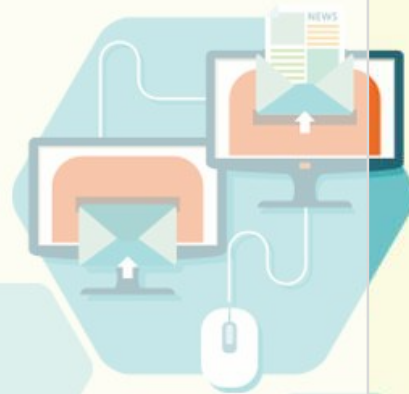
> library(MASS)

lda(formula, data, ...)

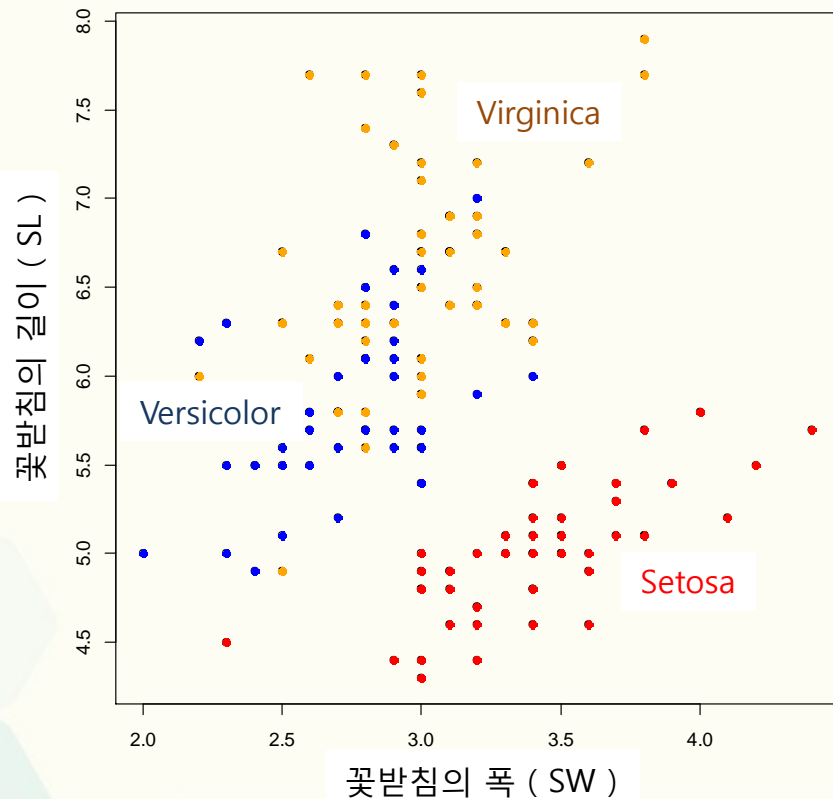
lda(x, grouping, ...)

기타 주요 전달인자 :

- prior : 사전확률, 고정값은 “propotional” (표본수 비례)
- Method : “moment”, “mle”, “mve”, “t” 고정값은 “moment”
- CV : leave-one-out 상호검증에 의한 “사후확률”제공
고정값은 FALSE



붓꽃 꽃받침의 폭과 길이



Prior

QDA : $Q_k(\mathbf{x}) = (\mathbf{x} - \mu_k)' \Sigma_k^{-1} (\mathbf{x} - \mu_k) + \log |\Sigma_k| - 2 \log f(y_{(k)})$ ↓

LDA : $\Sigma_k = \Sigma \quad k = 0, 1, \dots, K$

$$L_k = \mathbf{x}' \Sigma^{-1} \mu_k - (1/2) \mu_k' \Sigma^{-1} \mu_k + \log f(y_{(k)})$$
 ↑

- $f(y_{(k)}) = 1/(K + 1) \quad k = 0, 1, \dots, K$

- $f(y_{(k)}) = n_k/n \quad k = 0, 1, \dots, K$



2 그룹 판별의 경우

```
> newSP<-iris$SP  
> newSP[101:150]<-iris$SP[51:100]  
> newSP<- newSP[, drop=TRUE]
```

Versicolor , Virginica → “c”

Setosa → “s”

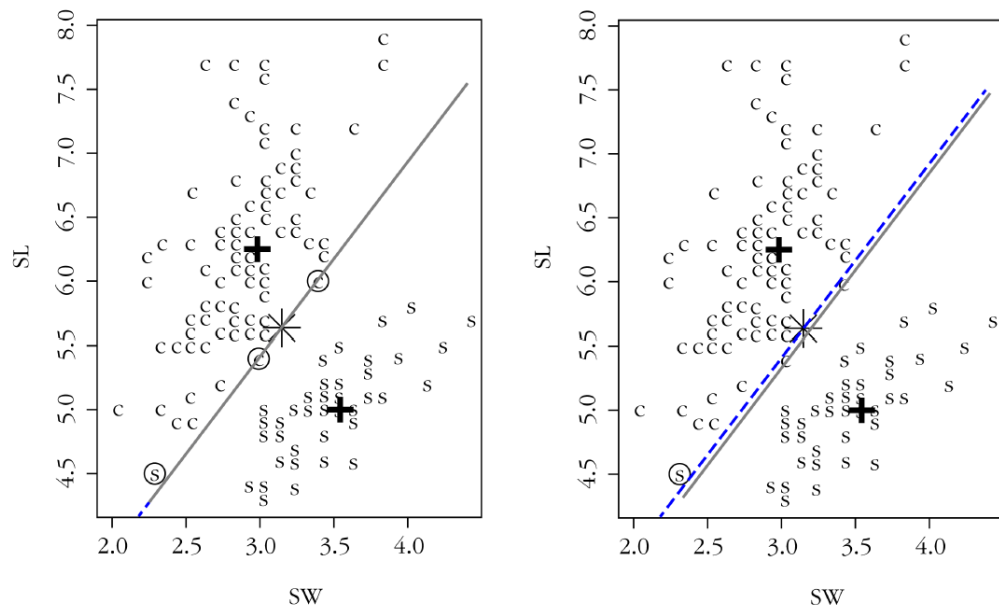
```
> library(MASS)  
> ( iris2a<- lda( newSP~SW+SL, prior=c(1,1)/2, data=iris ) )  
> ( iris2b<-lda( newSP~SW+SL, data=iris ) )
```

```
> par(mfrow=c(1,2)) # for black and white  
> plot.iris.lda(iris2a, c("s","c"), newSP, line=1, border=1)  
> plot.iris.lda(iris2b, c("s","c"), newSP, line=1, border=1)
```

plot.iris.lda() 함수는 부록에 수록되어 있음



Prior의 비교



[그림 7.1] 두 그룹 LDA 적용 예. (좌측은 사전확률 (0.5, 0.5), 우측은 사전확률 (1/3, 2/3). 잘못 분류된 결과들에 동그라미(O) 표시가 되어 있다.)



LDA, QDA & LRA

```
> ( iris3a<- lda( SP~SW+SL,data=iris) )      # LDA  
> iris3q<- qda( SP~SW+SL, data=iris)        # QDA  
> ( iris3m<- multinom( SP~SW+SL,data=iris) ) # LRA
```

```
> par(mfrow=c(1,3))
```

```
> plot.iris.lda(iris3a, border=1)  
> plot.iris.lda(iris3q, border=1)  
> plot.iris.mnom(iris3m, border=1)
```

plot.iris.mnom() 함수는 부록에 수록되어 있음



결과 : LDA & QDA

> lda(SP ~ SW + SL, data = iris)

Prior probabilities of groups:

st	vc	vg
0.3333333	0.3333333	0.3333333

Group means:

	SW	SL
st	3.428	5.006
vc	2.770	5.936
vg	2.974	6.588

Coefficients of linear discriminants:

	LD1	LD2
SW	2.768109	-2.0960764
SL	-2.141178	-0.8152721

Proportion of trace:

LD1	LD2
0.9628	0.0372

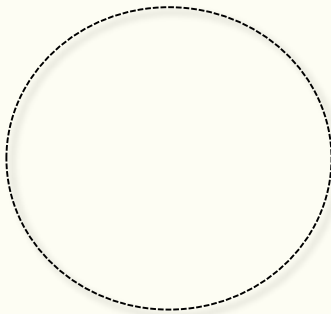
> qda(SP ~ SW + SL, data = iris)

Prior probabilities of groups:

st	vc	vg
0.3333333	0.3333333	0.3333333

Group means:

	SW	SL
st	3.428	5.006
vc	2.770	5.936
vg	2.974	6.588



결과 : LRA

```
> library(nnet)
> summary( multinom( SP ~ SW + SL, data = iris ) )
```

Coefficients:

	(Intercept)	SW	SL
vc	-92.09925	-40.58756	40.40327
vg	-105.10097	-40.18800	42.30095

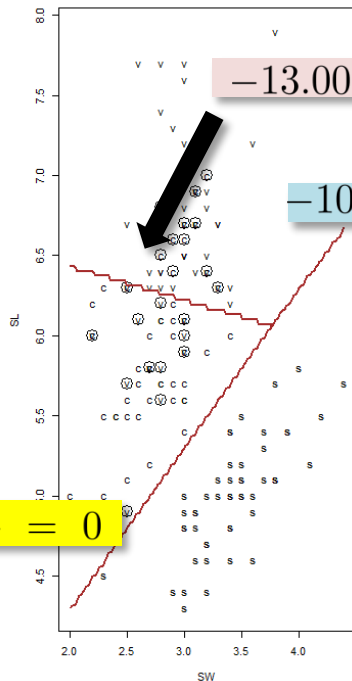
Std. Errors:

	(Intercept)	SW	SL
vc	26.27830	27.77771	9.142715
vg	26.37025	27.78874	9.131118

Residual Deviance: 110.425

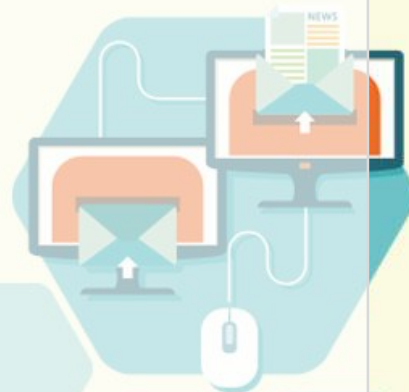
AIC: 122.425

$$-92.10 - 40.59 SW + 40.40 SL = 0$$

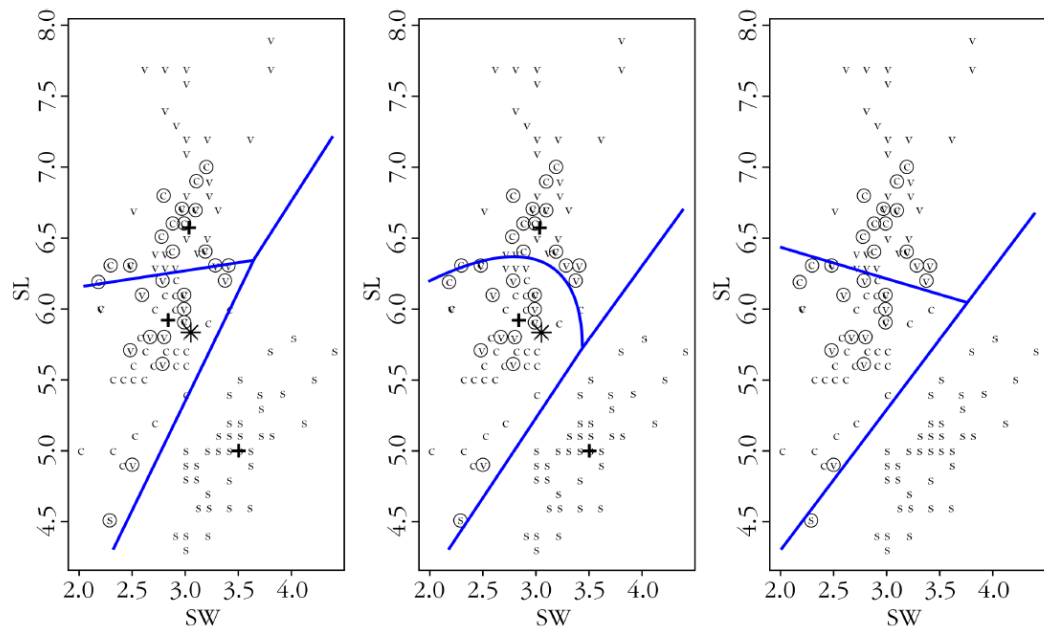


$$-13.00 + 0.40 SW + 1.90 SL = 0$$

$$-105.10 - 40.19 SW + 42.30 SL = 0$$



LDA, QDA & LRA



[그림 7.2] LDA, QDA, LRA 를 이용한 붓꽃자료 세 그룹 분류 결과
(좌측은 LDA, 가운데는 QDA, 우측은 LRA 잘못 분류된
결과들에 동그라미(O) 표시가 되어 있다.)



LDA, QDA & LRA : 2변수

```
> table(iris$SP, predict(iris3a)$class )  
> table(iris$SP, predict(iris3q)$class )  
> table(iris$SP, predict(iris3m) )
```

〈표 7.1〉 LDA, QDA, LRA 의 분류 결과 비교

	LDA			QDA			LRA		
	st	vc	cg	st	vc	vg	st	vc	vg
st	49	1	0	49	1	0	50	0	0
vc	0	36	14	0	37	13	0	38	12
vg	1	15	35	0	16	34	0	13	37



LDA, QDA & LRA : 4변수

```
> iris4<-lda(SP~. , data=iris)      # LDA  
> iris4q<-qda(SP~. , data=iris)     # QDA  
> iris4m<-multinom(SP~., data=iris) # Logistic  
  
> table(iris$SP, predict(iris4)$class )  
> table(iris$SP, predict(iris4q)$class )  
> table(iris$SP, predict(iris4m) )
```

〈표 7.2〉 네 변수를 이용한 붓꽃자료 LDA, QDA, LRA 적용 결과

	LDA			QDA			LRA		
	st	vc	vg	st	vc	vg	st	vc	vg
st	50	0	0	50	1	0	50	0	0
vc	0	48	2	0	48	2	0	49	1
vg	0	1	49	0	1	49	0	1	49



결과 : LDA (2) & LDA (4)

```
> lda(SP ~ SW + SL, data = iris)
```

Prior probabilities of groups:

st	vc	vg
0.3333333	0.3333333	0.3333333

Group means:

	SW	SL
st	3.428	5.006
vc	2.770	5.936
vg	2.974	6.588

Coefficients of linear discriminants:

	LD1	LD2
SW	2.768109	-2.0960764
SL	-2.141178	-0.8152721

Proportion of trace:

LD1	LD2
0.9628	0.0372

```
> lda(SP ~ ., data = iris)
```

Prior probabilities of groups:

st	vc	vg
0.3333333	0.3333333	0.3333333

Group means:

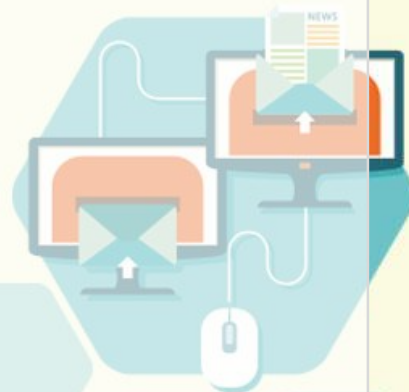
	SL	SW	PL	PW
st	5.006	3.428	1.462	0.246
vc	5.936	2.770	4.260	1.326
vg	6.588	2.974	5.552	2.026

Coefficients of linear discriminants:

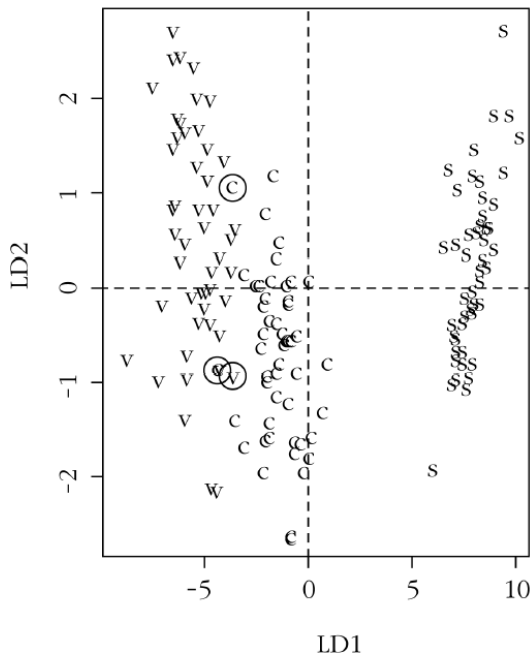
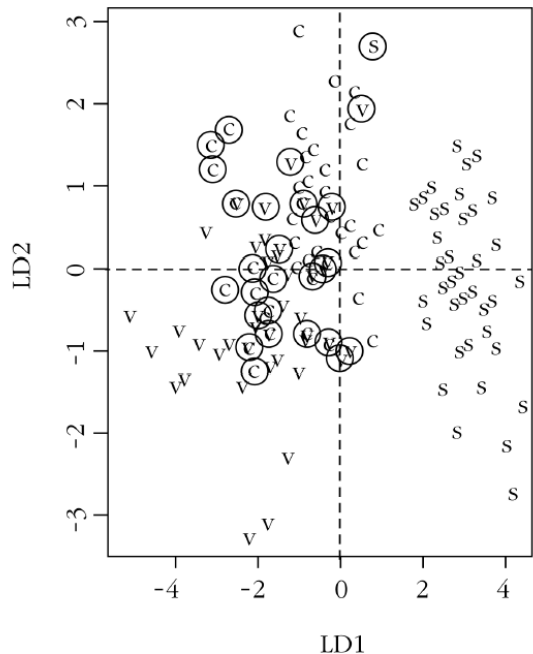
	LD1	LD2
SL	0.8293776	0.02410215
SW	1.5344731	2.16452123
PL	-2.2012117	-0.93192121
PW	-2.8104603	2.83918785

Proportion of trace:

LD1	LD2
0.9912	0.0088



LDA : 2변수, 4변수



[그림 7.4] 붓꽃자료에 대한 LDA 결과 비교
(좌측 (SW, SL) 사용, 우측 (SW, SL, PW, PL) 사용 경우
잘못 분류된 결과들에 동그라미(O) 표시가 되어 있다)

```
> par(mfrow=c(1,2))
```

```
> plot.iris.discriminant(iris3a)
```

```
> plot.iris.discriminant(iris4)
```

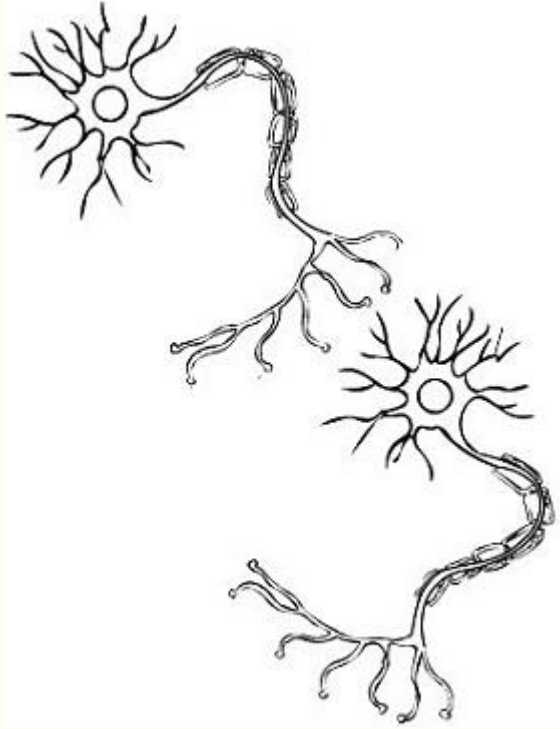
plot.iris.discriminant() 함수는
부록에 수록되어 있음



③ 신경망 분류방법



neuron & triggering



LRA의 그림표현 : 4변수

> multinom(SP ~ . , data = iris)

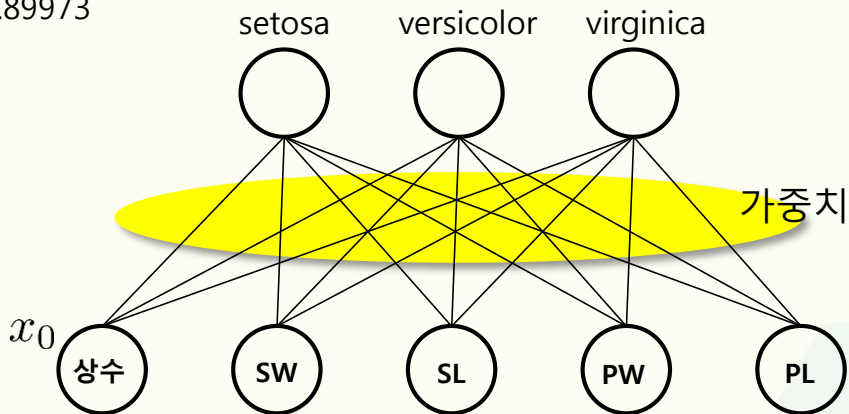
Coefficients:

	(Intercept)	SL	SW	PL	PW
st	0	0	0	0	0
vc	18.69	-5.46	-8.70	14.24	-3.10
vg	-23.84	-7.92	-15.37	23.66	15.14

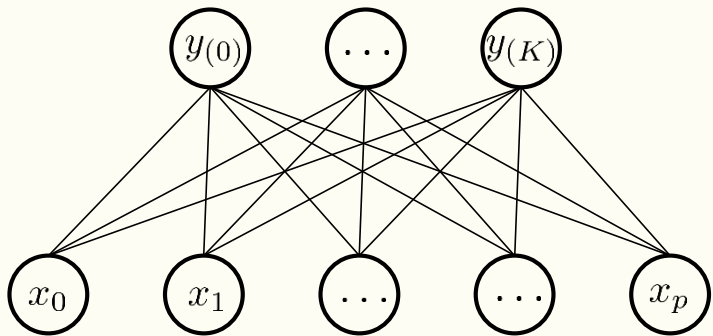
가중치 (weight : Wts)

Residual Deviance: 11.89973

AIC: 31.89973



LRA의 수리적 표현



$$y = (y_{(0)}, y_{(1)}, \dots, y_{(K)})' \sim \text{Multinom}(1, p_0, p_1, \dots, p_K)$$

$$x = (1, x_1, \dots, x_p)' \quad \beta_k = (\beta_{k0}, \beta_{k1}, \dots, \beta_{kp})'$$

$$p_k = \frac{\exp(\beta'_k x)}{\sum_{k=0}^K \exp(\beta'_k x)} \quad k = 0, 1, \dots, K$$

Softmax 함수



은닉변수층이 있는 모형의 예

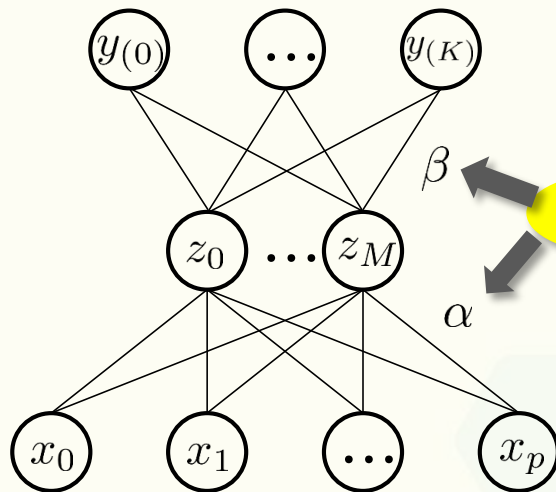
$$y = (y_{(0)}, y_{(1)}, \dots, y_{(K)})' \sim \text{Multinom}(1, p_0, p_1, \dots, p_K)$$

$$z_m \sim \text{Bin}(1, q_m) \quad m = 0, 1, \dots, M$$

$$p_k = \frac{\exp(\beta'_k q)}{\sum_{k=0}^K \exp(\beta'_k q)} \quad k = 0, 1, \dots, K$$

$$q_m = \frac{1}{1 + \exp(\alpha'_m x)} \quad m = 0, 1, \dots, M$$

Sigmoid 함수



가중치



모형적합 방법

$$R(\alpha, \beta) + \lambda J(\alpha, \beta)$$



$$y_i \in \{y_{(0)}, \dots, y_{(K)}\} \quad x_{i\cdot} = (x_{i0}, \dots, x_{ip})' \quad i = 1, \dots, n$$

$$y_{i(k)} = I(y_i = y_{(k)})$$

적합결여도 :

$$R(\alpha, \beta) = \sum_{i=1}^n \sum_{k=0}^K (y_i - f_k(x_{i\cdot}))^2$$

잔차제곱합(회귀목적)

$$R(\alpha, \beta) = - \sum_{i=1}^n \sum_{k=0}^K y_{i(k)} \log p_k(x_{i\cdot})$$

엔트로피(분류목적)

거칠기 벌칙 : (모형의 단순성 추구)

$$J(\alpha, \beta) = \sum_{k,m} \beta_{km}^2 + \sum_{mi} \alpha_{mi}^2$$

제곱합벌칙

$$J(\alpha, \beta) = \sum_{k,m} \frac{\beta_{km}^2}{1 + \beta_{km}^2} + \sum_{mi} \frac{\alpha_{mi}^2}{1 + \alpha_{mi}^2}$$

소거벌칙



④ 신경망 분류방법의 적용 예



R의 nnet 함수

```
> library(nnet)
```

```
nnet(formula, data, ...)
```

```
nnet(x, y,...)
```

기타 주요 전달인자 :

- size : 히든노드의 수
- Wts : 추정해야 할 각 weight 의 초기값.
- rang : 초기 weight 값이 없는 경우 배정할 확률난수 범위
- decay : 모형적합에서 사용할 거칠기 벌칙항에 대한 가중값
- maxit : 최적화 과정의 최대 허용 반복 횟수
- weights : 자료의 가중치



class.ind & max.col

```
> ( smp<- iris$SP[sample(1:150,9)] )
```

```
[1] vg st vg vc vc vg vg st st
```

```
Levels: st vc vg
```

```
> as.integer(smp)
```

```
[1] 3 1 3 2 2 3 3 1 1
```

```
> class.ind(smp)
```

```
      st vc vg
```

```
[1,] 0 0 1
```

```
[2,] 1 0 0
```

```
[3,] 0 0 1
```

```
[4,] 0 1 0
```

```
...
```

```
[9,] 1 0 0
```

```
> max.col(class.ind(smp))
```

```
[1] 3 1 3 2 2 3 3 1 1
```



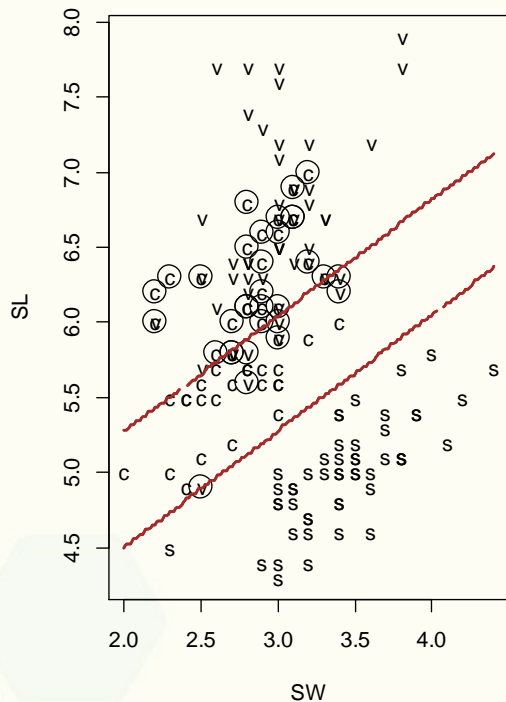
신경망

```
> irx <- iris[,-5]
> spc <- class.ind( iris$SP )
> irx3n1 <- nnet(spc~ SW+SL , data=irx, size = 1, rang = 0.1, decay = 5e-4, maxit = 300 )
> irx3n2 <- nnet(spc~ SW+SL , data=irx, size = 2, rang = 0.1, decay = 5e-4, maxit = 300 )
# weights: 15
initial value 113.276176
iter 10 value 84.446863
iter 240 value 37.693508
final value 37.692786
converged

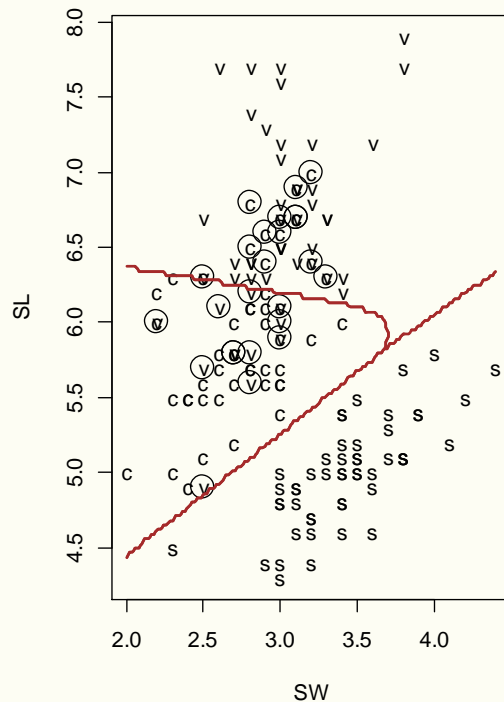
> par(mfrow=c(1,2))
> plot.iris.mnom(irx3n1)
> plot.iris.mnom(irx3n2)
```



히든노드의 수에 따른 신경망 적합결과



히든노드 수 1개



히든노드 수 2개

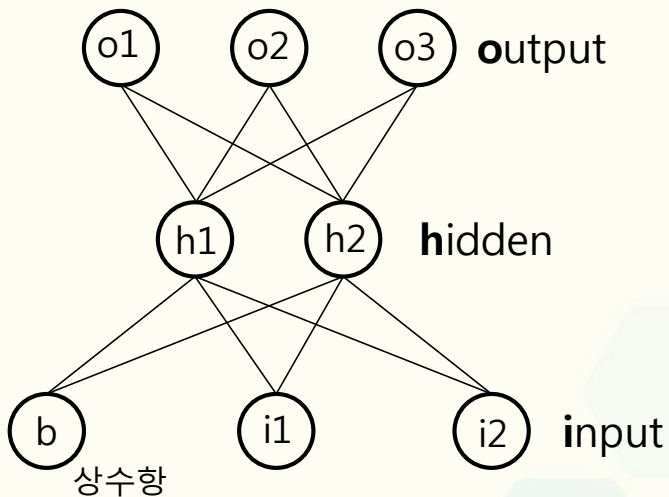


가중치의 추정

> summary(irx3n2)

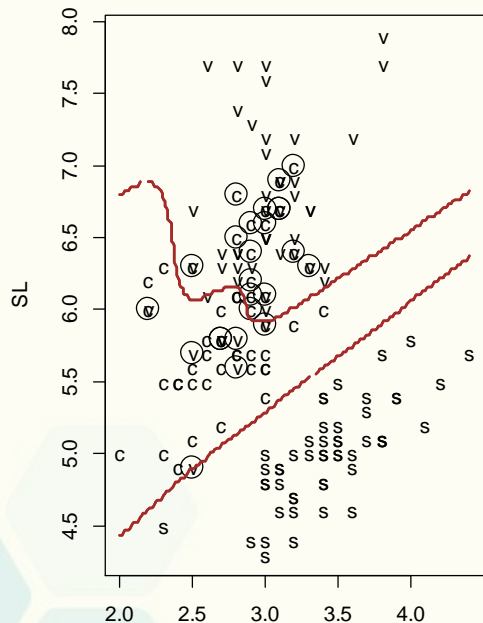
a 2-2-3 network with 15 weights
options were - decay=5e-04

b->h1	i1->h1	i2->h1
24.77	6.25	-8.41
b->h2	i1->h2	i2->h2
6.93	-0.18	-1.15
b->o1	h1->o1	h2->o1
-4.05	13.11	-3.04
b->o2	h1->o2	h2->o2
-2.85	-11.26	8.82
b->o3	h1->o3	h2->o3
2.73	-6.21	-8.26

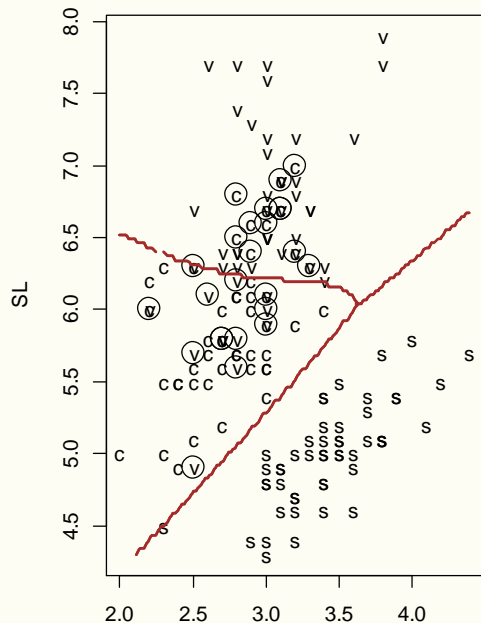


과적합과 벌칙항

```
> irx3n5 <- nnet(spc~ SW+SL , data=irx, size = 5, rang = 0.1, decay = 5e-4, maxit = 300 )  
> irx3n6 <- nnet(spc~ SW+SL , data=irx, size = 5, rang = 0.1, decay = 0.01, maxit = 300 )
```



sw decay=0.0005



sw decay=0.01

히든노드 수 5개

$$R(\alpha, \beta) + \lambda J(\alpha, \beta)$$

decay





다음시간 안내

분류분석 (2)

