Module 18 Multivariate Analysis for Genetic data Session 11 Discriminant Analysis I

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Discriminant analysis: Aims

- Group separation
- Dimension reduction: from p variables to k discriminators with k < p.
- Classification of new cases

Discriminant Analysis: the data matrix

Ind.	X_1	<i>X</i> ₂		X_p	Group
1	X ₁₁	X ₁₂		X_{1p}	1
2	X_{21}	X_{22}		X_{2p}	1
:	:	:	:	:	:
n.	Υ.	Υ .		Υ .	1
$\frac{n_1}{1}$	<i>X</i> _{n11}	<i>X</i> _{n12}		X_{n_1p}	
1	X ₁₁	X_{12}		X_{1p}	2
2	X_{21}	X_{22}	• • •	X_{2p}	2
:					
n_2	X_{n_21}	$X_{n_2 2}$	• • •	X_{n_2p}	2
1	X ₁₁	X ₁₂		X_{1p}	m
2	X_{21}	X_{22}	• • •	X_{2p}	m
•		•		•	
•		•	•		
n _m	X_{n_m1}	X_{n_m2}		X_{n_mp}	m

Introduction

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- Given various biochemical measurements, is this person healthy or diseased?
- Given the variables of this wheat kernel, to which of the known varieties does it belong?
- ...
- One can distinguish between two-group and multiple group problems.

Two-group linear discriminant analysis

Criteria for designing a classification rule:

- small probability of misclassification
- take prevalence into account (prior probabilities)
- take the cost of misclassification into account

Two-group linear discriminant analysis

Some basic definitions:

- \bullet π_1 and π_2 represent population 1 and 2.
- $f_1(x)$ and $f_2(x)$ represent the multivariate probability densities for each population.
- $\Omega = R_1 \cup R_2$ is the partitioned sample space for outcome x.
- If x falls in R₁, the case is classified as π₁, else in π₂.
- p_1 is the prior probability of pertaining to π_1 , p_2 the prior probability of pertaining to π_2 (prevalence)
- Misclassification probabilities:
 - **1** $P(2|1) = P(\mathbf{X} \in R_2|\pi_1) = \int_{R_2} f_1(\mathbf{x}) d\mathbf{x}$
 - 2 $P(1|2) = P(\mathbf{X} \in R_1|\pi_2) = \int_{R_1} f_2(\mathbf{x}) d\mathbf{x}$

Cost matrix

		Predicted class		
		π_1	π_2	
True	π_1	0	c(2 1)	
Class	π_2	c(1 2)	0	

c(1|2) and c(2|1) are not necessarily equal

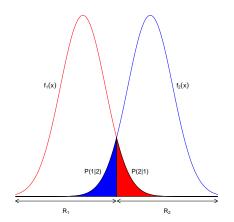
ECM = Expected Cost of Misclassification

$$P ext{ (from } \pi_1 \cap \text{ classified } \pi_2) = P (2|1) \cdot p_1$$

$$P ext{ (from } \pi_2 \cap \text{ classified } \pi_1) = P (1|2) \cdot p_2$$

$$ECM = c(1|2)P(1|2)p_2 + c(2|1)P(2|1)p_1$$

Classification rule: minimizing ECM



ECM Rule

Introduction

The regions that minimize the ECM are:

$$R_1: rac{f_1(\mathbf{x})}{f_2(\mathbf{x})} \geq 1$$
 $R_2: rac{f_1(\mathbf{x})}{f_2(\mathbf{x})} < 1$

If there is differential prevalence:

$$R_1: \frac{f_1(\mathbf{x})}{f_2(\mathbf{x})} \ge \frac{p_2}{p_1} \qquad R_2: \frac{f_1(\mathbf{x})}{f_2(\mathbf{x})} < \frac{p_2}{p_1}$$

If there is differential cost:

$$R_1: \frac{f_1(\mathbf{x})}{f_2(\mathbf{x})} \ge \frac{c(1|2)}{c(2|1)}$$
 $R_2: \frac{f_1(\mathbf{x})}{f_2(\mathbf{x})} < \frac{c(1|2)}{c(2|1)}$

And if we have both differential prevalence and differential cost:

$$R_1: \frac{f_1(\mathbf{x})}{f_2(\mathbf{x})} \ge \frac{c(1|2)}{c(2|1)} \cdot \frac{p_2}{p_1} \qquad R_2: \frac{f_1(\mathbf{x})}{f_2(\mathbf{x})} < \frac{c(1|2)}{c(2|1)} \cdot \frac{p_2}{p_1}$$

Introduction

Two normal populations with equal covariance matrices

For continuous X, we assume multivariate normality:

$$f_1(\mathbf{x}) = \frac{1}{(2\pi)^{p/2} |\mathbf{\Sigma}|^{\frac{1}{2}}} e^{-\frac{1}{2}(\mathbf{x} - \mu_1)'\mathbf{\Sigma}^{-1}(\mathbf{x} - \mu_1)}$$

$$f_2(\mathbf{x}) = rac{1}{(2\pi)^{p/2} |\mathbf{\Sigma}|^{rac{1}{2}}} e^{-rac{1}{2}(\mathbf{x}-\mu_2)'\mathbf{\Sigma}^{-1}(\mathbf{x}-\mu_2)}$$

Introduction

Error rate

Two-group linear discriminant analysis

Sample based ECM Rule: assign observation \mathbf{x} to population 1 if

$$(\bar{\mathbf{x}}_1 - \bar{\mathbf{x}}_2)'\mathbf{S}_{\rho}^{-1}\mathbf{x} - \frac{1}{2}(\bar{\mathbf{x}}_1 - \bar{\mathbf{x}}_2)'\mathbf{S}_{\rho}^{-1}(\bar{\mathbf{x}}_1 + \bar{\mathbf{x}}_2) \geq \ln\left(\left(\frac{c(1|2)}{c(2|1)}\right)\left(\frac{\rho_2}{\rho_1}\right)\right)$$

where S_p is the pooled covariance matrix:

$$\mathbf{S}_p = \frac{n_1 - 1}{n_1 + n_2 - 2} \mathbf{S}_1 + \frac{n_2 - 1}{n_1 + n_2 - 2} \mathbf{S}_2$$

Two-group linear discriminant analysis

Define:

$$\mathbf{a} = \mathbf{S}_n^{-1}(\mathbf{\bar{x}}_1 - \mathbf{\bar{x}}_2) \qquad y = \mathbf{a}'\mathbf{x}$$

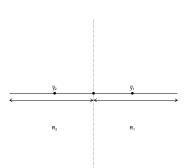
Note that:

$$y_i = \mathbf{a}' \mathbf{x}_i \qquad \overline{y}_1 = \mathbf{a}' \overline{\mathbf{x}}_1 \qquad \overline{y}_2 = \mathbf{a}' \overline{\mathbf{x}}_2$$

With equals costs and priors, the ECM rule for R_1 boils down to the univariate rule:

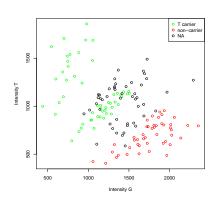
$$y_i > \frac{1}{2}(\overline{y}_1 + \overline{y}_2)$$

y is the classifier or linear discriminant function.



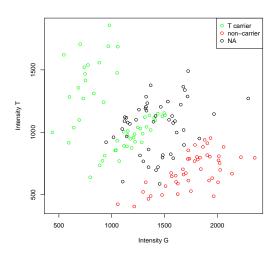
Example: SNP intensities and called genotypes

	SNP	iG	iΤ
1	TT	641	1037
2	GT	1207	957
3	TT	1058	1686
4	GG	1348	466
5	GT	1176	948
6	GG	1906	912
:			:
12	NA	947	920
:	:	:	:



- Calling algorithm assigns missings to "difficult" genotypes
- Could we reasonably predict if these are carriers of the T allele?

Re-plotting

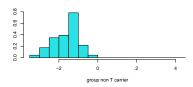


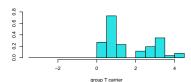
Two-group LDA output

Model: Carrier status \sim iT + iG

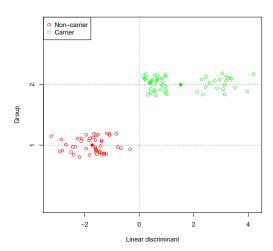
		group	means
	Prior	iT	iG
non T carrier	0.47	691.59	1758.78
T carrier	0.53	1133.56	1037.44

Linear discriminant function:





Graphical representation





Prediction of missings

			Posterior	prob.
	Prediction	LD1	non.T.carrier	T.carrier
12	T carrier	1.25	0.01	0.99
20	non T carrier	-0.25	0.59	0.41
21	non T carrier	-0.98	0.94	0.06
27	T carrier	1.15	0.02	0.98
28	T carrier	0.22	0.24	0.76
29	non T carrier	-1.83	1.00	0.00
	:	:	:	:

Error rate

Two-group QDA

Introduction

Under the assumption of multivariate normality with $\Sigma_1 \neq \Sigma_2$, using the same ECM principle, a quadratic classification rule is obtained.

Sample based ECM Rule: assign observation x to population 1 if

$$-\frac{1}{2}\mathbf{x}'(\mathbf{S}_1^{-1}-\mathbf{S}_2^{-1})\mathbf{x}+(\overline{\mathbf{x}}_1'\mathbf{S}_1^{-1}-\overline{\mathbf{x}}_2'\mathbf{S}_2^{-1})\mathbf{x}-k\geq \ln\left(\left(\frac{c(1|2)}{c(2|1)}\right)\left(\frac{\rho_2}{\rho_1}\right)\right)$$

with

$$k = \frac{1}{2} \ln \left(\frac{|\mathbf{S}_1|}{|\mathbf{S}_2|} \right) + \frac{1}{2} (\overline{\mathbf{x}}_1' \mathbf{S}_1^{-1} \overline{\mathbf{x}}_1 - \overline{\mathbf{x}}_2' \mathbf{S}_2^{-1} \overline{\mathbf{x}}_2)$$

Sample covariance matrices

	Non-carriers					
	iG	iT				
iG	71677.24	24891.89				
iΤ	24891.89	20914.78				

	Carriers	
	iG	iT
iG	77553.35	-23117.55
iT	-23117.55	81695.66

Predictions for missings

	LDA			QDA			
	Prediction	LD1	non.T.carrier	T.carrier	Prediction	non.T.carrier.1	T.carrier.1
12	T carrier	1.25	0.01	0.99	T carrier	0.00	1.00
20	non T carrier	-0.25	0.59	0.41	T carrier	0.00	1.00
21	non T carrier	-0.98	0.94	0.06	non T carrier	0.79	0.21
27	T carrier	1.15	0.02	0.98	T carrier	0.00	1.00
28	T carrier	0.22	0.24	0.76	T carrier	0.00	1.00
29	non T carrier	-1.83	1.00	0.00	non T carrier	0.99	0.01
32	T carrier	0.10	0.31	0.69	T carrier	0.00	1.00
35	non T carrier	-0.64	0.83	0.17	non T carrier	0.54	0.46
41	T carrier	0.87	0.04	0.96	T carrier	0.00	1.00
47	T carrier	0.83	0.04	0.96	T carrier	0.00	1.00
48	T carrier	0.99	0.02	0.98	T carrier	0.00	1.00
52	T carrier	0.04	0.36	0.64	T carrier	0.03	0.97
58	non T carrier	-0.95	0.93	0.07	non T carrier	0.84	0.16
62	non T carrier	-0.52	0.78	0.22	T carrier	0.09	0.91
65	non T carrier	-0.80	0.90	0.10	non T carrier	0.69	0.31
69	non T carrier	-0.71	0.87	0.13	non T carrier	0.74	0.26
72	non T carrier	-0.70	0.86	0.14	non T carrier	0.71	0.29
75	non T carrier	-0.67	0.85	0.15	T carrier	0.17	0.83
76	T carrier	1.24	0.01	0.99	T carrier	0.00	1.00
80	non T carrier	-0.18	0.53	0.47	T carrier	0.13	0.87
81	T carrier	0.44	0.13	0.87	T carrier	0.00	1.00
83	T carrier	-0.08	0.45	0.55	T carrier	0.00	1.00
87	T carrier	-0.01	0.40	0.60	T carrier	0.23	0.77
89	T carrier	0.35	0.17	0.83	T carrier	0.00	1.00
92	T carrier	1.04	0.02	0.98	T carrier	0.00	1.00
95	non T carrier	-1.28	0.98	0.02	non T carrier	0.93	0.07
101	T carrier	1.07	0.02	0.98	T carrier	0.00	1.00
102	T carrier	0.89	0.03	0.97	T carrier	0.00	1.00
104	T carrier	0.96	0.03	0.97	T carrier	0.00	1.00
106	T carrier	1.18	0.01	0.99	T carrier	0.00	1.00
108	non T carrier	-1.09	0.96	0.04	non T carrier	0.92	0.08
110	T carrier	1.05	0.02	0.98	T carrier	0.00	1.00
115	T carrier	0.40	0.15	0.85	T carrier	0.00	1.00
118	non T carrier	-1.19	0.97	0.03	non T carrier	0.68	0.32
121	T carrier	1.16	0.01	0.99	T carrier	0.00	1.00
122	T carrier	-0.08	0.46	0.54	T carrier	0.03	0.97
123	non T carrier	-0.59	0.82	0.18	T carrier	0.45	0.55
126	T carrier	0.34	0.18	0.82	T carrier	0.00	1.00
127	T carrier	0.79	0.05	0.95	T carrier	0.00	1.00
128	T carrier	0.85	0.04	0.96	T carrier	0.00	1.00
129	T carrier	-0.10	0.47	0.53	T carrier	0.00	1.00
131	T carrier	0.40	0.15	0.85	T carrier	0.00	1.00
134	T carrier	-0.05	0.43	0.57	T carrier	0.00	1.00
135	T carrier	-0.06	0.44	0.56	T carrier	0.00	1.00
138	T carrier	1.16	0.01	0.99	T carrier	0.00	1.00
139	T carrier	0.76	0.05	0.95	T carrier	0.00	1.00
144	non T carrier	-0.44	0.73	0.27	T carrier	0.44	0.56
145	non T carrier	-0.23	0.58	0.42	T carrier	0.00	1.00

Error rates and Confusion matrix

- It is of interest to evaluate the performance of a classification rule.
- There are several criteria to do so.
- Actual error rate (AER, density dependent)

$$\mathsf{AER} = p_1 \int_{\hat{R}_2} f_1(\mathbf{x}) d\mathbf{x} + p_2 \int_{\hat{R}_1} f_2(\mathbf{x}) d\mathbf{x}$$

 Apparent error rate (APER, not density dependent) based on the confusion matrix

		Predicted class		
		π_1	π_2	
True	π_1	n_{11}	n ₁₂	
Class	π_2	n_{21}	n ₂₂	

APER obtained as

$$APER = \frac{n_{12} + n_{21}}{n_1 + n_2}$$

APER underestimates the AER

References

- Hand, D.J. (1981) Discrimination and Classification. Wiley, New York.
- Johnson & Wichern, (2002) Applied Multivariate Statistical Analysis, 5th edition, Prentice Hall, Chapter 11.
- Lachenbruch, P.A. (1975) Discriminant Analysis. Hafner Press, New York.