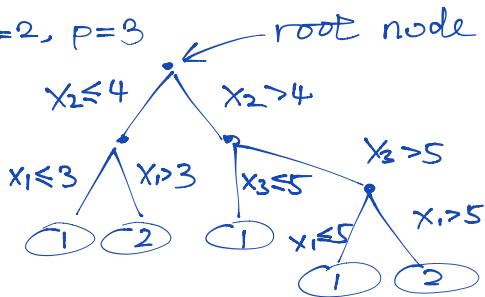


Classification Tree

Want to model $\theta_j(\underline{x}) = P(G=j | \underline{X}=\underline{x})$ for $j=1, \dots, k$

$k=2, p=3$



Question: How to construct (grow) tree?

Recursive partitioning

At a given node, define $B = \{(g_i, \underline{x}_i) : i \in I(B)\}$
 \uparrow
 $1, \dots, k$

For $\underline{x}_i \in B$, we have

$$\hat{\theta}_j(B) = \frac{1}{n(B)} \sum_{i \in I(B)} I(g_i = j) = \frac{n_j(B)}{n(B)}$$

 \uparrow
 $\#$ of observations in B
 \uparrow
 $=$ proportions of group j in node B .

Now define possible new nodes: B_1, B_2 with $B = B_1 \cup B_2$

$$\hat{\theta}_j(B_1) = \frac{1}{n(B_1)} \sum_{i \in I(B_1)} I(g_i = j) = \frac{n_j(B_1)}{n(B_1)} \quad (j=1, \dots, k) \quad \text{disjoint}$$

$$\hat{\theta}_j(B_2) = \frac{1}{n(B_2)} \sum_{i \in I(B_2)} I(g_i = j) = \frac{n_j(B_2)}{n(B_2)}$$

Define $D(B_1, B_2; B) = \sum_{j=1}^k \left\{ n_j(B_1) \ln \left(\frac{n_j(B_1)}{n(B_1)} \right) + n_j(B_2) \ln \left(\frac{n_j(B_2)}{n(B_2)} \right) - n_j(B) \ln \left(\frac{n_j(B)}{n(B)} \right) \right\}$
 $\uparrow \quad \uparrow \quad \uparrow$
variables fixed

- Find B_1, B_2 s.t. $B_1 \cup B_2 = B$ to maximize D .
 \uparrow
many choices

- restrict maximization to simple one variable splits

$$B_1 = \{(g_i, \underline{x}_i) \in B, x_{i,l} \leq d\} \quad \leftarrow \text{threshold}$$

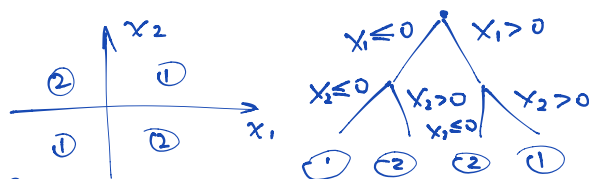
$$B_2 = \{(g_i, \underline{x}_i) \in B, x_{i,l} > d\}$$

\uparrow
 $l=1, \dots, p$

- also require other constraints e.g. $n(B_1), n(B_2) \geq 5$.

Does procedure always work?

Example: $k=2, p=2$



How does tree algorithm work here?

- $P(G=j | X_1) = \frac{1}{2}$ (for $j=1, 2$)
 $P(G=j | X_2) = \frac{1}{2}$

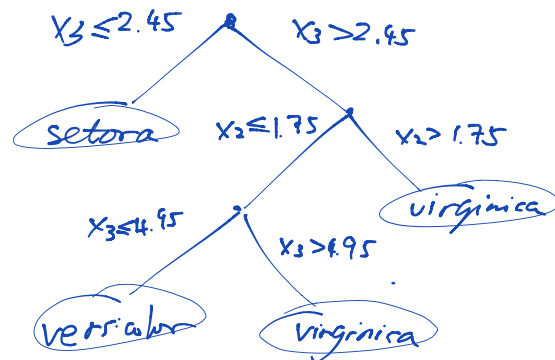
- recursive partitioning algorithm has trouble getting started

- better performance if allow splitting along linear combinations (i.e. projections) of variables.

Example Iris data

3 species { setosa, virginica, versicolor }
 4 vars { x_1 = sepal length, x_2 = sepal width, x_3 = petal length, x_4 = petal width }

- compare favourably to LDA
- error rate = 4/150 (tree)
- error rate = 3/150 (LDA)



Regression models for multivariate data

- repeated measures

Multivariate Analysis of Variance (MANOVA)

Problem k treatments (or groups)

n_i subjects in treatment i

Multivariate response

p vectors $\vec{x}_{ij} \begin{cases} i=1, \dots, k \\ j=1, \dots, n_i \end{cases}$

Model: $\vec{x}_{ij} = \mu_i + \epsilon_{ij}$ $i=1, \dots, k, j=1, \dots, n_i$

where $\{\epsilon_{ij}\}$ are independent $N_p(0, C)$ random vectors i.e. $\vec{x}_{ij} \sim N_p(\mu_i, C)$

One question of interest: Is there a difference between k treatments? (For example, is $\mu_1 = \mu_2 = \dots = \mu_k$?)

Univariate ($p=1$) case: Decompose total sum of squares

$$SS_{\text{Total}} = \sum_{i=1}^k \sum_{j=1}^{n_i} (x_{ij} - \bar{x})^2 = \sum_{i=1}^k n_i (\bar{x}_i - \bar{x})^2 + \sum_{i=1}^k \sum_{j=1}^{n_i} (x_{ij} - \bar{x}_i)^2$$

\downarrow overall sample mean \downarrow sample mean for treatments

$$SS_{\text{Total}} = SS_{\text{between group}} + SS_{\text{within group}}$$

To test $H_0: \mu_1 = \dots = \mu_k$, we compare SS_{between} to SS_{within}

Test statistic $F = \frac{SS_{\text{between}}(k-1)}{SS_{\text{within}}(n-k)} \sim F_{k-1, n-k}$ under H_0

$$SS_{\text{Total}} = \sum_{i=1}^k \sum_{j=1}^{n_i} (\vec{x}_{ij} - \bar{\vec{x}})(\vec{x}_{ij} - \bar{\vec{x}})^T = \underbrace{\sum_{i=1}^k n_i (\bar{\vec{x}}_i - \bar{\vec{x}})(\bar{\vec{x}}_i - \bar{\vec{x}})^T}_{SS_{\text{between}}} + \underbrace{\sum_{i=1}^k \sum_{j=1}^{n_i} (\vec{x}_{ij} - \bar{\vec{x}}_i)(\vec{x}_{ij} - \bar{\vec{x}}_i)^T}_{SS_{\text{within}}}$$

Question:

How to compare SS_{between} to SS_{within}