

$$g(x_1, x_2) = 10.5 \mathbb{1}_{x_1 > 5} + 55 \mathbb{1}_{x_1 \leq 5} \mathbb{1}_{x_2 \leq 5} + 10.5 \mathbb{1}_{x_1 \leq 5} \mathbb{1}_{x_2 > 5}$$

Binary Tree Models are linear models where the variables are carved into partial spaces like categorical variables.

where all categorical variables are binarized already

One regression tree algorithm (there are many)

- ① Begin with $\mathcal{D} = \langle X, \hat{y} \rangle, X = [\vec{x}_1, \vec{x}_2, \dots, \vec{x}_p] = \begin{bmatrix} \vec{x}_1 & \vec{x}_2 & \dots & \vec{x}_p \\ x_{11} & x_{12} & \dots & x_{1p} \\ x_{21} & x_{22} & \dots & x_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ x_{n1} & x_{n2} & \dots & x_{np} \end{bmatrix}$
- ② Consider all possible orthogonal-to-axis splits i.e.

$x_1 \leq x_{(1)1}, x_1 \leq x_{(2)1}, \dots, x_1 \leq x_{(n-1)1},$
 $x_2 \leq x_{(1)2}, x_2 \leq x_{(2)2}, \dots, x_2 \leq x_{(n-1)2},$
 \vdots
 $x_p \leq x_{(1)p}, x_p \leq x_{(2)p}, \dots, x_p \leq x_{(n-1)p} \Rightarrow (n-1)p \text{ possible splits}$

Sorted $\vec{x}_{\cdot 1}$ values from min-max.

For each split there are two putative daughter nodes.

For each split we assign \hat{y} = \bar{y} of the observations that landed in the node. Calculate SSE in both daughters.

- ③ Locate the split rule with smallest #-obs-weighted average SSE:

$$SSE_w := \frac{n_L SSE_L + n_R SSE_R}{n_L + n_R}$$

n_L := # of observations in left daughter node

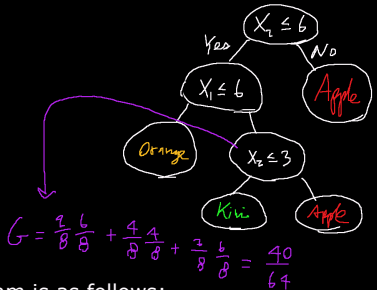
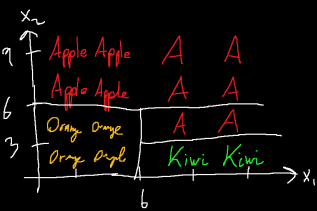
n_R := # of observations in right daughter node

This algorithm is "greedy" because at each juncture it takes the locally optimal solution and these algorithms never find global optimums.

- ④ Create the optimal split and the two daughter nodes.
- ⑤ Repeat steps 2-4 on the daughter nodes unless daughter node has N_0 observations (e.g. $N_0 = 5$) and unless there's only one unique y .

Let $y = \{C_1, C_2, \dots, C_k\}$ classification problem

Classification Trees



$$G = \frac{2}{8} \frac{6}{8} + \frac{4}{8} \frac{4}{8} + \frac{2}{8} \frac{6}{8} = \frac{40}{64}$$

One classification tree algorithm is as follows:

- ① Same as Regression Tree Algorithm above
- ② Consider all splits as in the Regression Tree Algorithm. For all daughter nodes set \hat{y} = Mode[y's in the daughter node] and calculate a heterogeneity metric called the "Gini"

$$G := \sum_{l=1}^K \hat{p}_l (1 - \hat{p}_l) \text{ where } \hat{p}_l := \frac{\# \{y_i \text{ in category } l\}}{\# \text{ obs. in node}}$$

- ③ Same as Regression Tree Algorithm above. Take the #-obs-weighted Gini average:

$$G_w := \frac{n_L G_L + n_R G_R}{n_L + n_R}$$

- ④ Same as Regression Tree Algorithm above.
- ⑤ Same as Regression Tree Algorithm above. Stop at default $N_0 = 1$ or if node is completely homogeneous.