

Decision trees

Readings for today

- Chapter 8: Tree-based methods. James, G., Witten, D., Hastie, T., & Tibshirani, R. (2013). An introduction to statistical learning: with applications in R (Vol. 6). New York: Springer

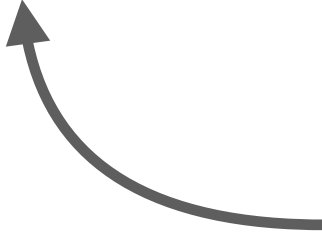
Topics

1. Decision trees
2. Classification trees
3. Trees vs. linear models

Decision trees

Interpretability

Linear Models: $\hat{y}_i = \hat{f}(x_i) = \hat{\beta}_0 + \hat{\beta}_1 x_i$



unit change in y that happens with each
unit change in x

Assumes: 1. $Y \sim N(\mu, \sigma)$
2. X is *iid*

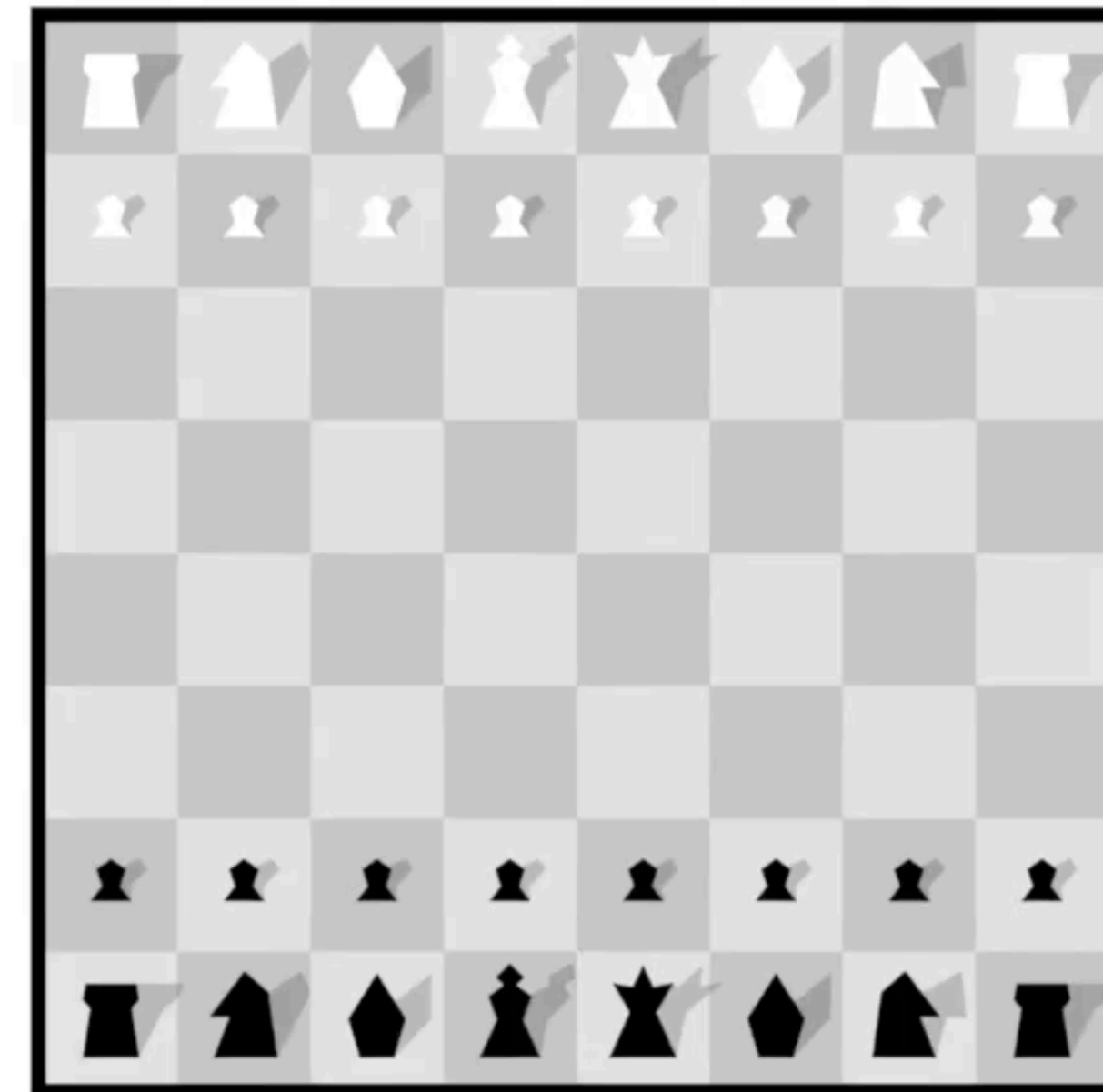
As soon as your data violates these assumptions, the interpretability of your effects is impaired.

Tree methods

Goal: Treat $f(X)$ as a series of “if/then” rules.

Approach: Identify the best decision rules for parsing X to explain Y .

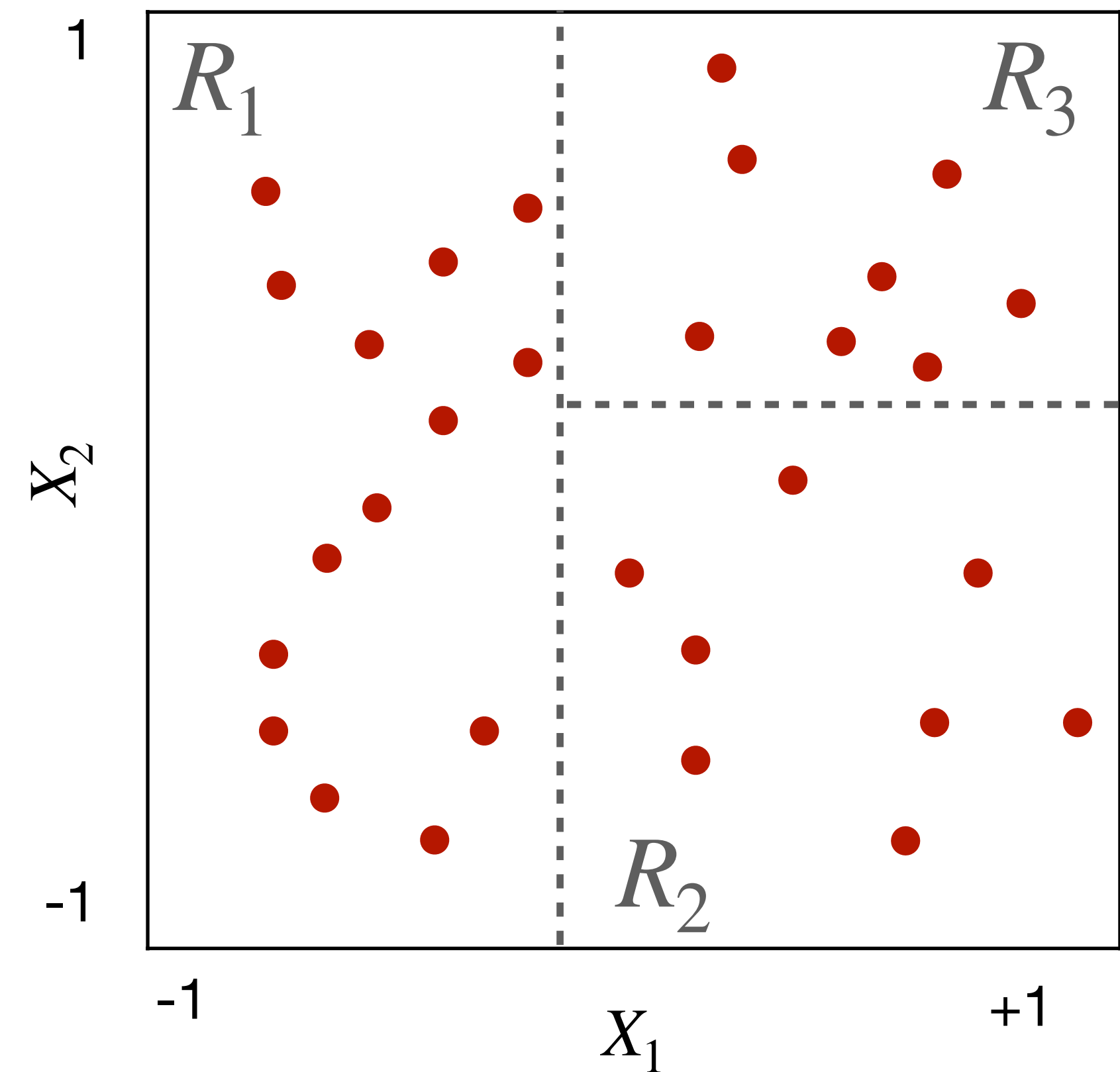
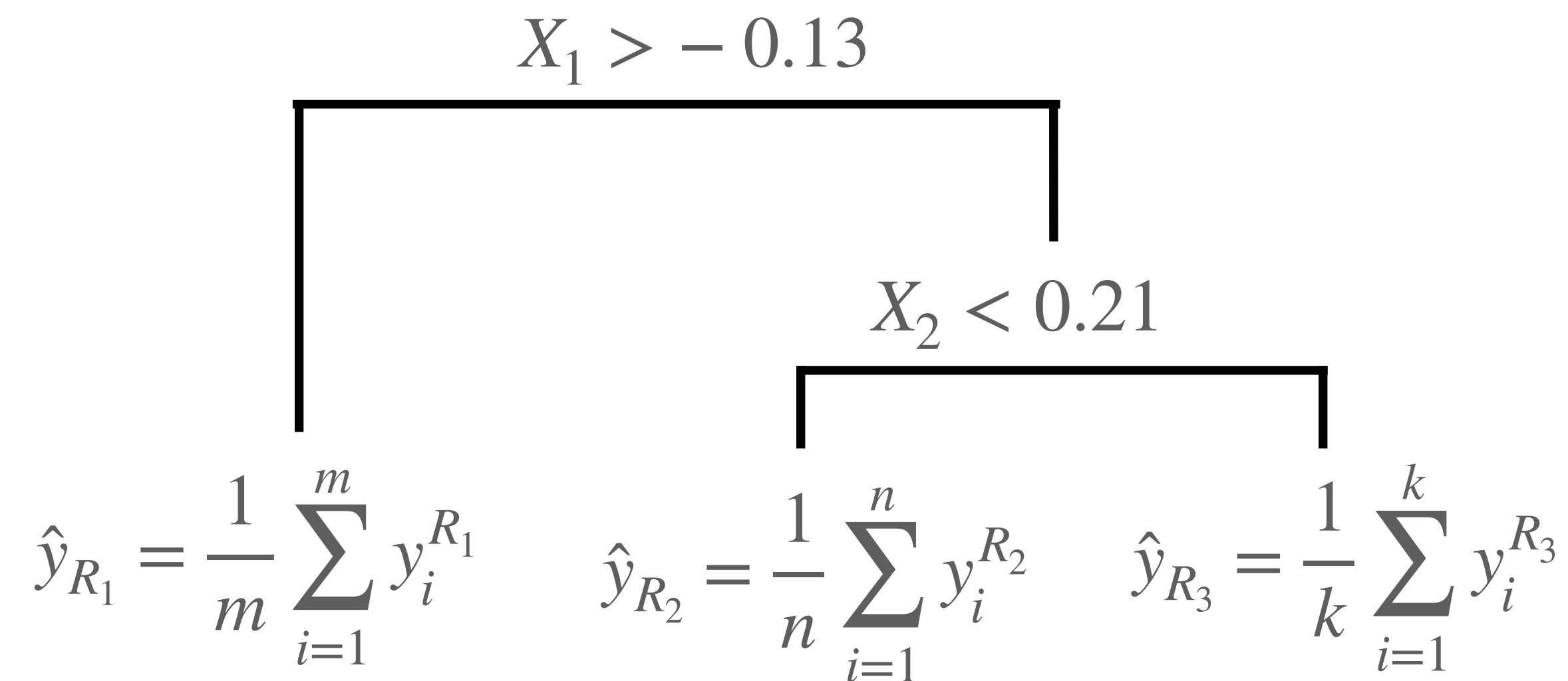
Non-parametric way of identifying subregions in X relevant to Y .



Google DeepMind

Recursive Binary Splitting

- Steps: 1. Divide X recursively into j subregions R_1, \dots, R_j .
2. Assign \hat{y} as the mode of every y_i in region R_m



Objective function

Goal: $\min(\sum_{j=1}^J \sum_{i \in R_j}^n (y_i - \hat{y}_{R_j})^2) \rightarrow RSS$

\hat{y}_{R_j} is the mean/modal value of all observations in region R_j .

Top-down: Split most relevant feature first and then work your way down successively.

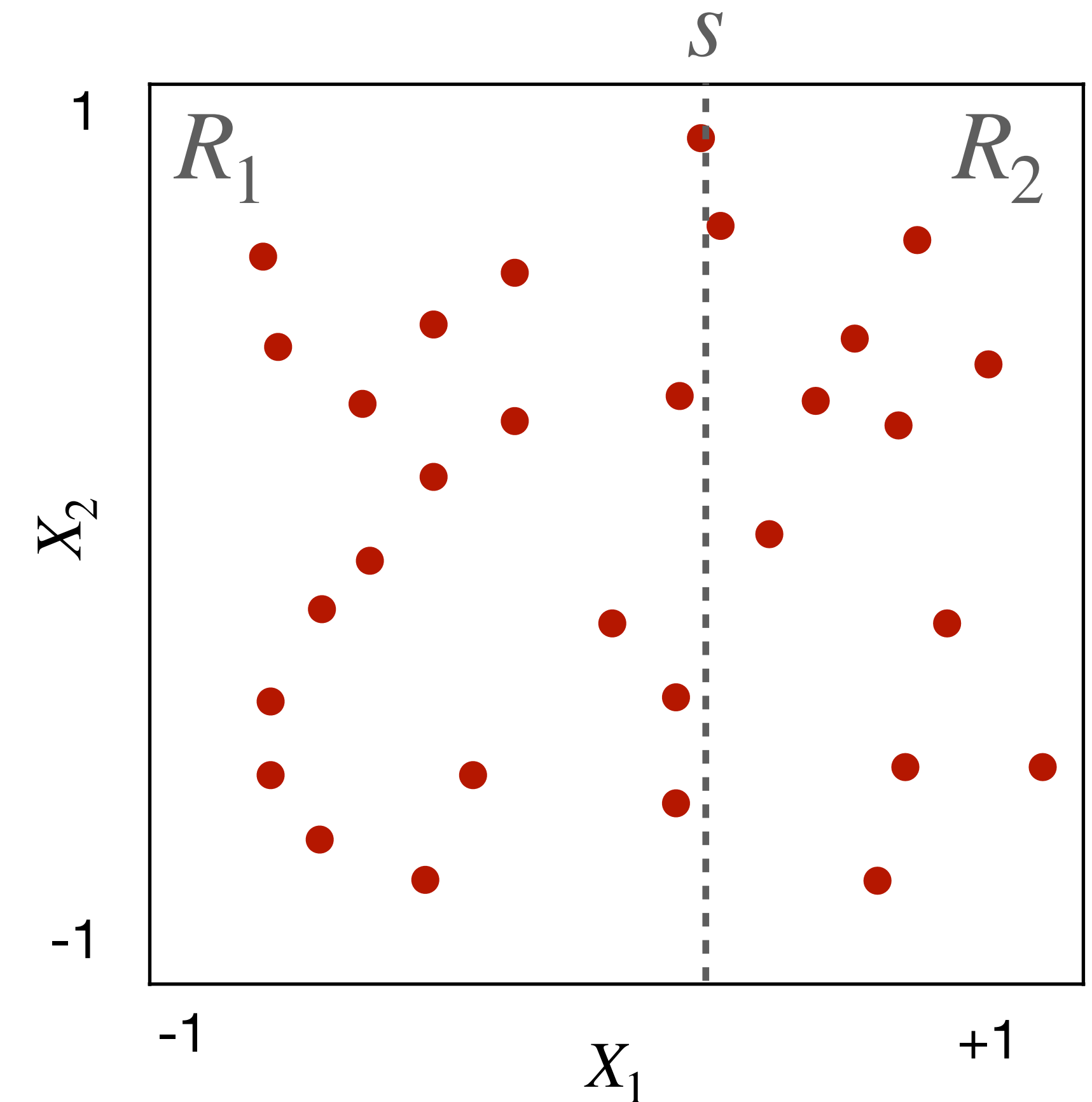
Greedy: *Only* the “best” split (i.e., split that explains most variance) is chosen at each step.

Splitting

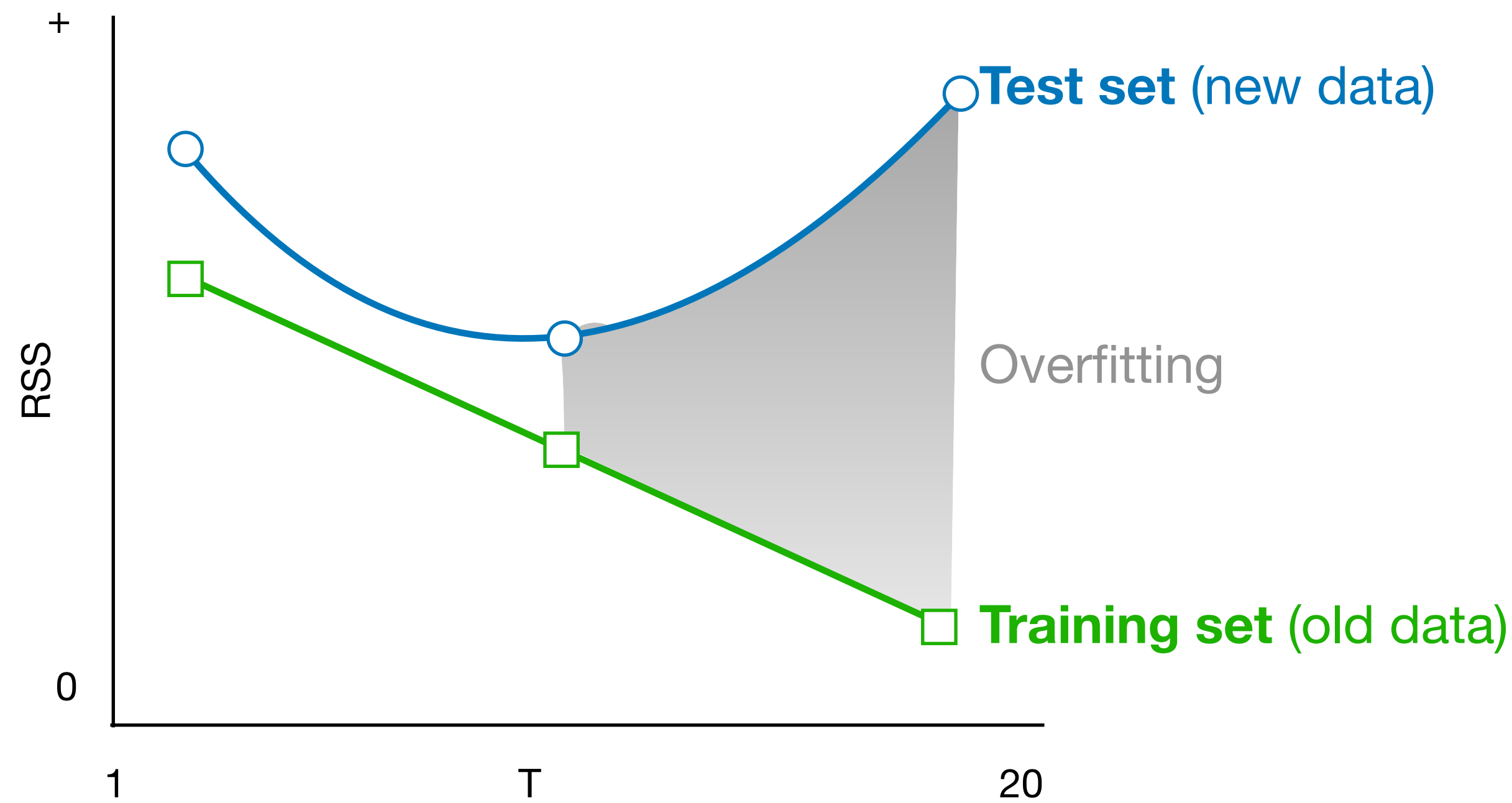
Choosing the right split means finding the best border, s , that minimizes two residual estimates.

$$\begin{array}{cc} \text{Region 1} & \text{Region 2} \\ \hline R_1(j, s) = \{X \mid X_j < s\} & R_2(j, s) = \{X \mid X_j \geq s\} \end{array}$$

minimize: $\sum_{i: x_i \in R_1(j, s)} (y_i - \hat{y}_{R_1})^2 + \sum_{i: x_i \in R_2(j, s)} (y_i - \hat{y}_{R_2})^2$



Bias-variance tradeoff



Number of regions you choose to split along, T , determines the flexibility of your model

$\uparrow T = \uparrow \text{variance}$

Cost-Complexity Tuning:

$$\min \left(\sum_{m=1}^T \sum_{i: x_i \in R_m}^n (y_i - \hat{y}_{R_m})^2 + \hat{\alpha} |T| \right)$$

$\uparrow \alpha = \downarrow T$

sparsity parameter to be tuned

Classification trees

Classification trees

Goal: Create a tree for qualitative (categorical) Y .

classification error rate $E_m = 1 - \max_k (\hat{p}_{mk})$ Proportion of training observations in region m that are members of class k .

Objective: $\min(\sum_{j=1}^J \sum_{i \in m}^K E_m)$ Minimize the classification error across classes.

Node purity

Node purity: Measure of uniqueness of categories in each R_m .

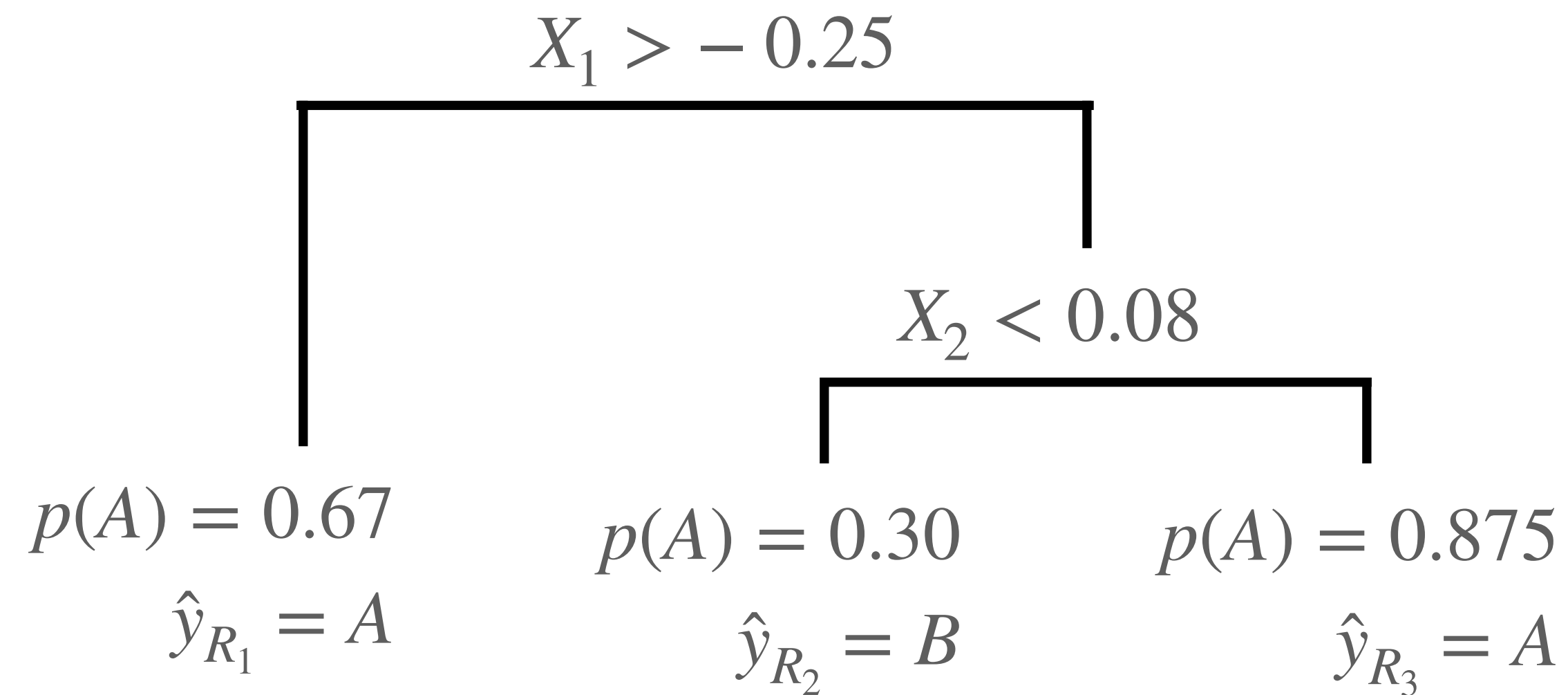
1. Gini Index (G): $G = \sum_{k=1}^K \hat{p}_{mk}(1 - \hat{p}_{mk})$

2. Cross-entropy (D): $D = \sum_{k=1}^K \hat{p}_{mk} \log(\hat{p}_{mk})$

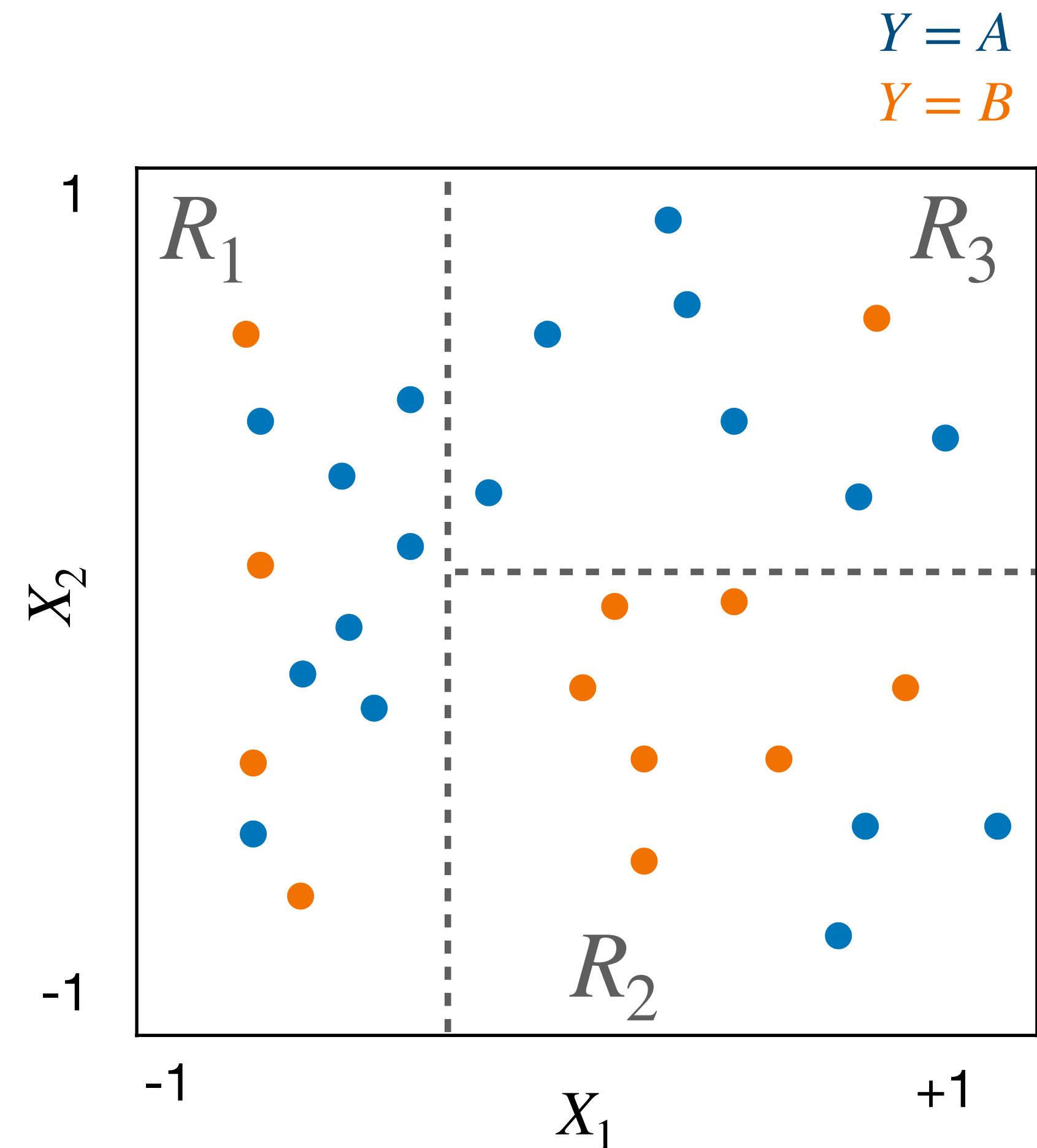
- $\downarrow G = \uparrow$ purity
- $\downarrow D = \uparrow$ purity

Both measures determine whether a node/region contains predominantly members of one class.

Example: 2 group classification

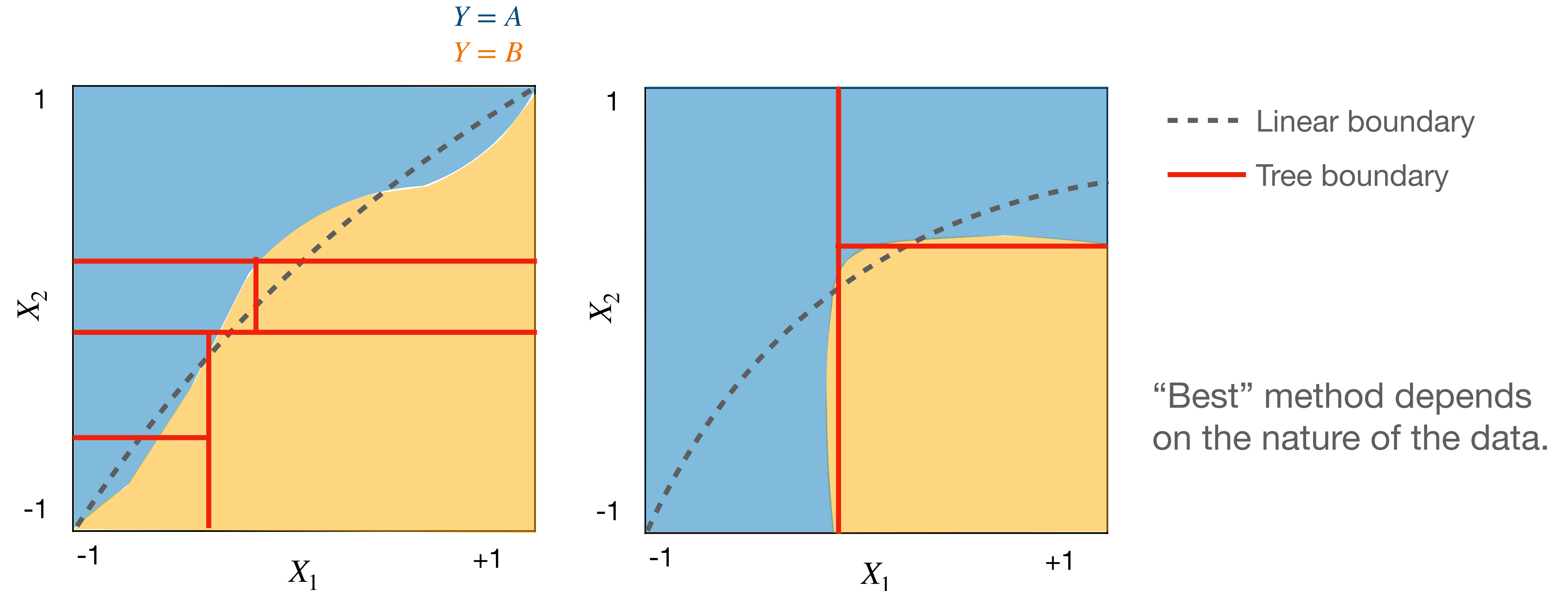


$G = 0.54$ ← Moderate node purity.



Trees vs. linear models

Trees vs. linear models



The pros and cons

Advantages:

- Easy to explain
- Easily visualized
- No need for dummy coding
- “Mirrors” human heuristics
- Applicable to data that does not meet the assumptions of linear or parametric models.

Disadvantages:

- Lower predictive accuracy.
- Non-robust



small changes in data have huge impacts on model fits.

Take home message

- Decision trees are intuitive, but finicky, methods for determining $X \rightarrow Y$ relationships that are especially powerful in contexts where the data does not meet the assumptions of standard linear models.