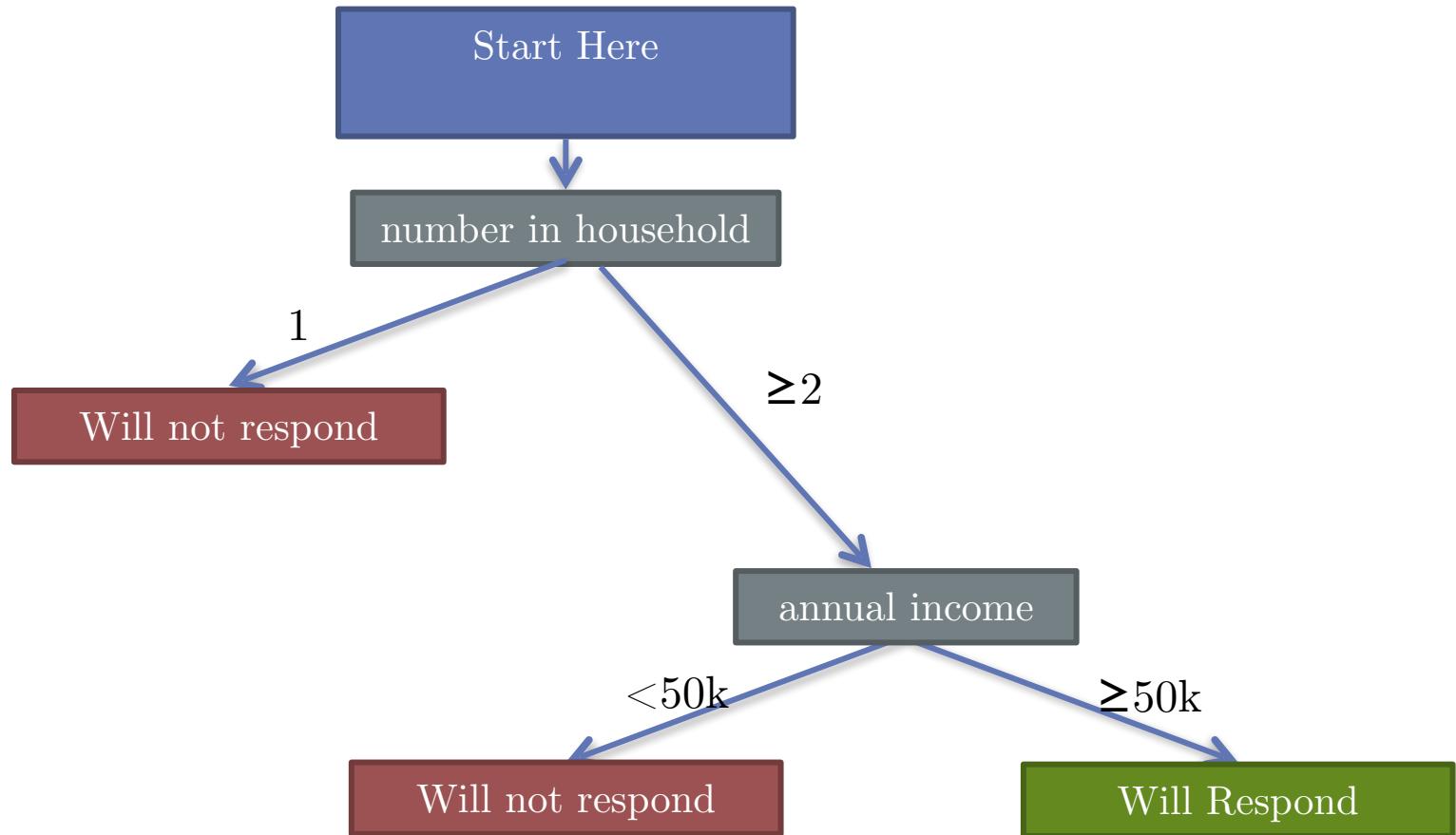


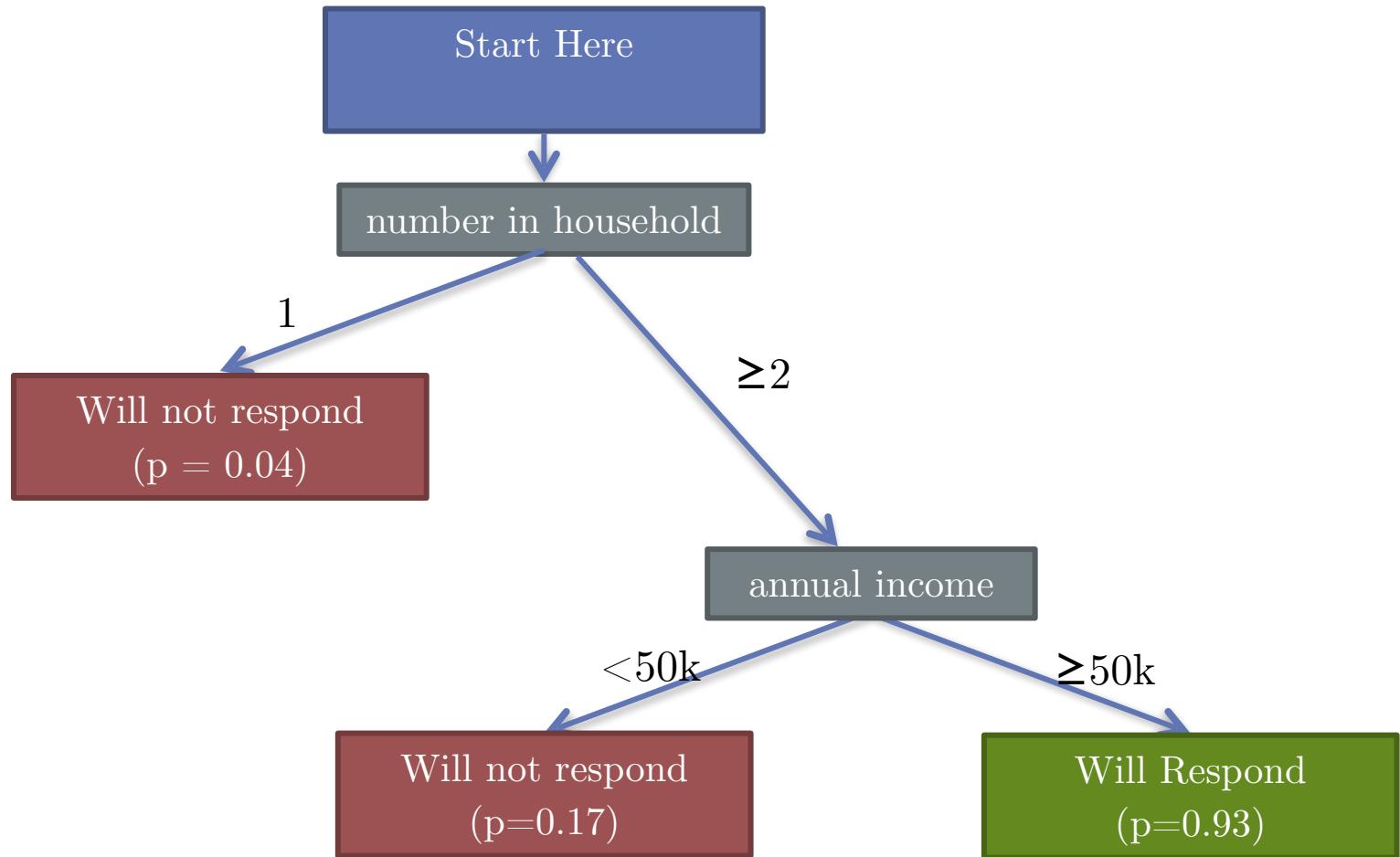
Classification And Regression Trees (CARTs)

a.k.a. Decision Trees

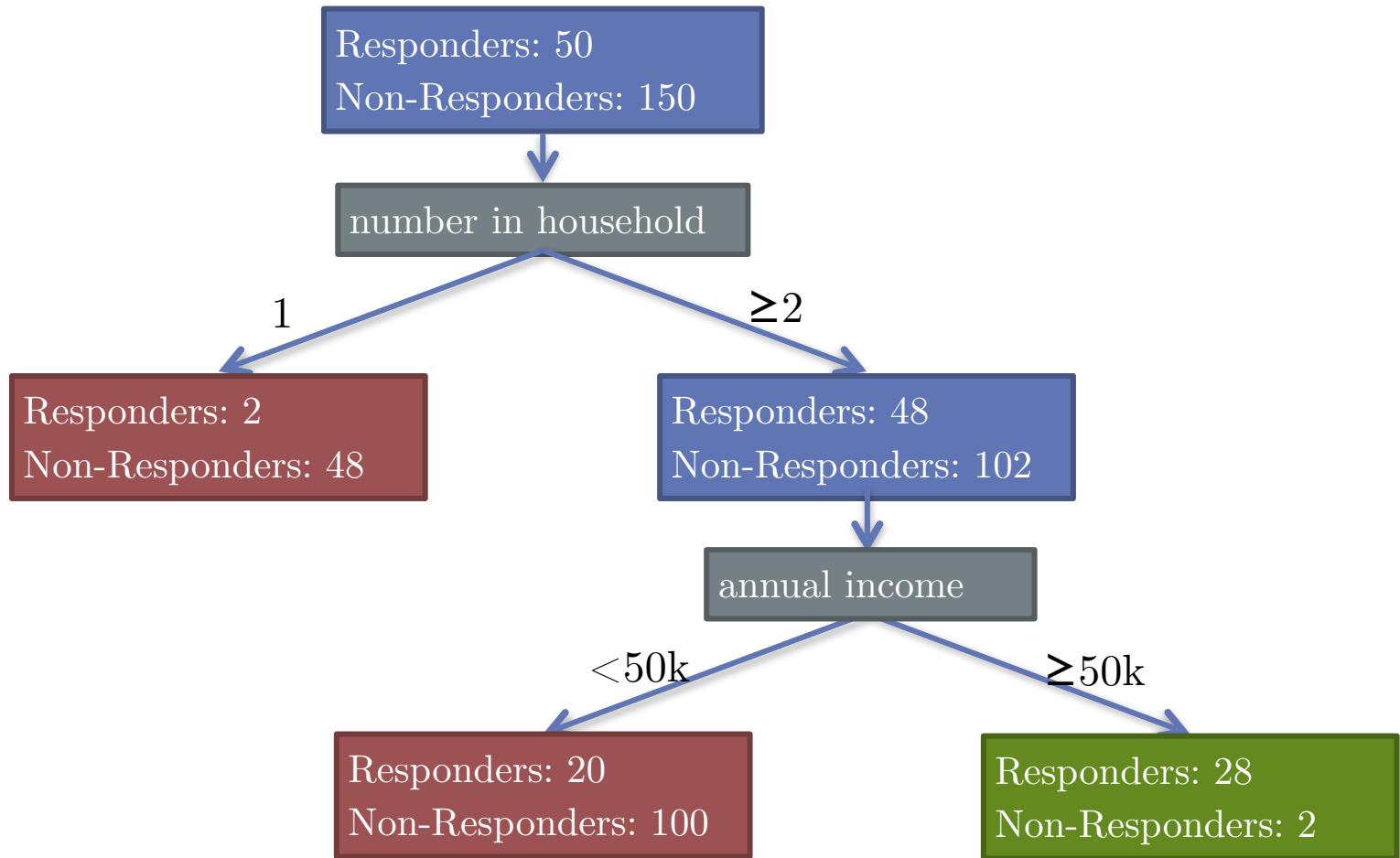
A Decision Tree Model



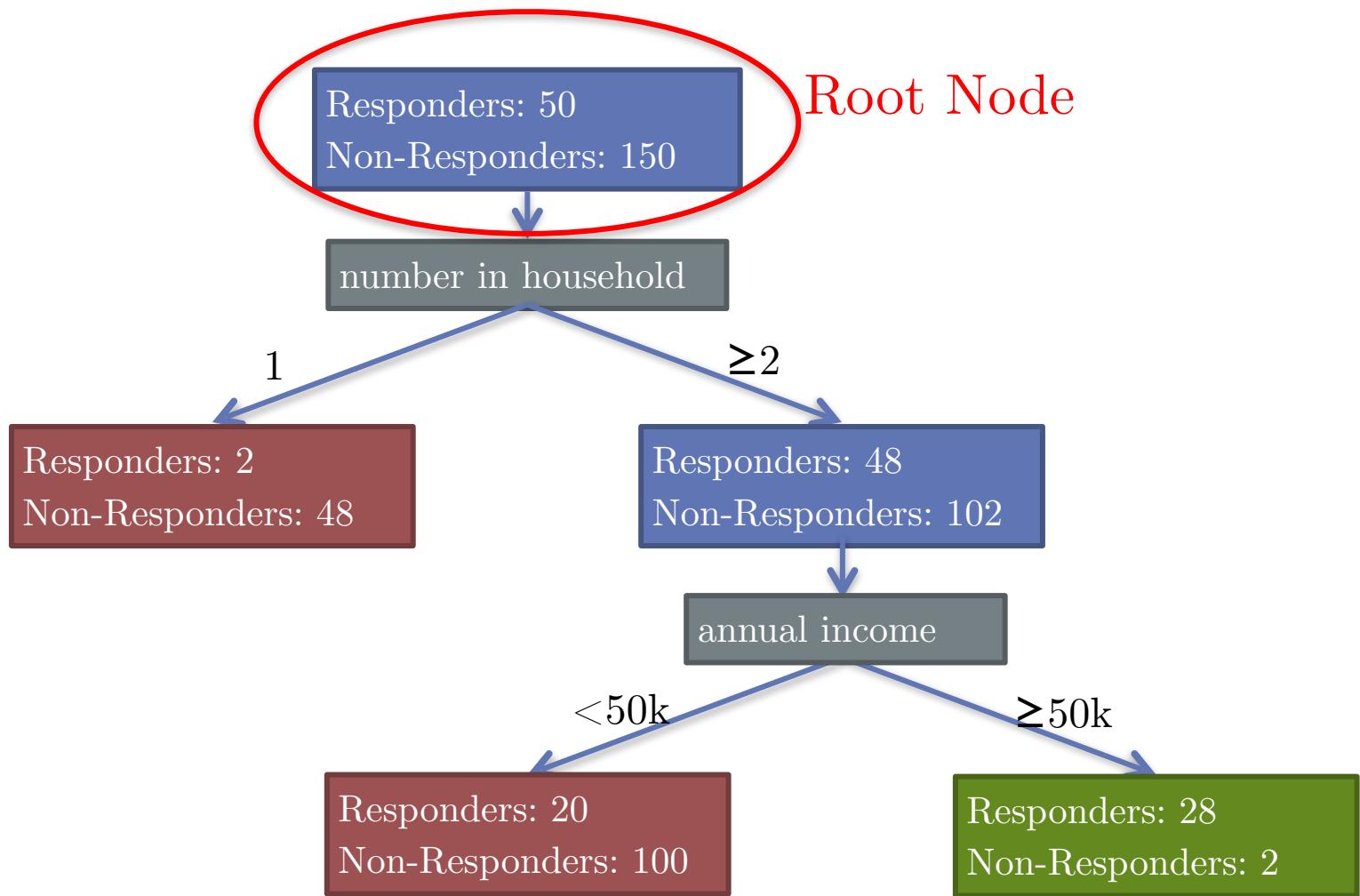
A Decision Tree Model



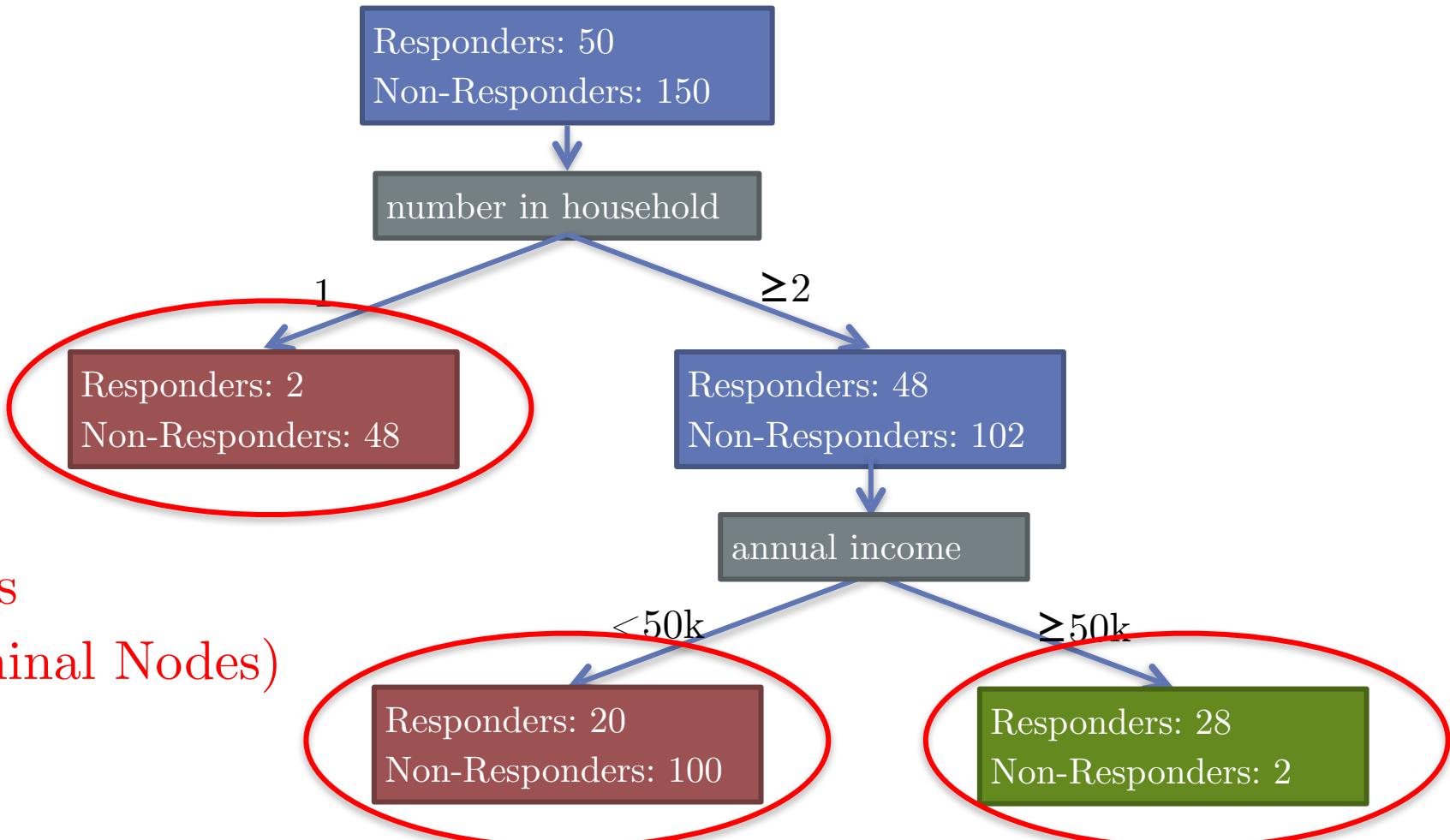
Decision Tree Model Creation



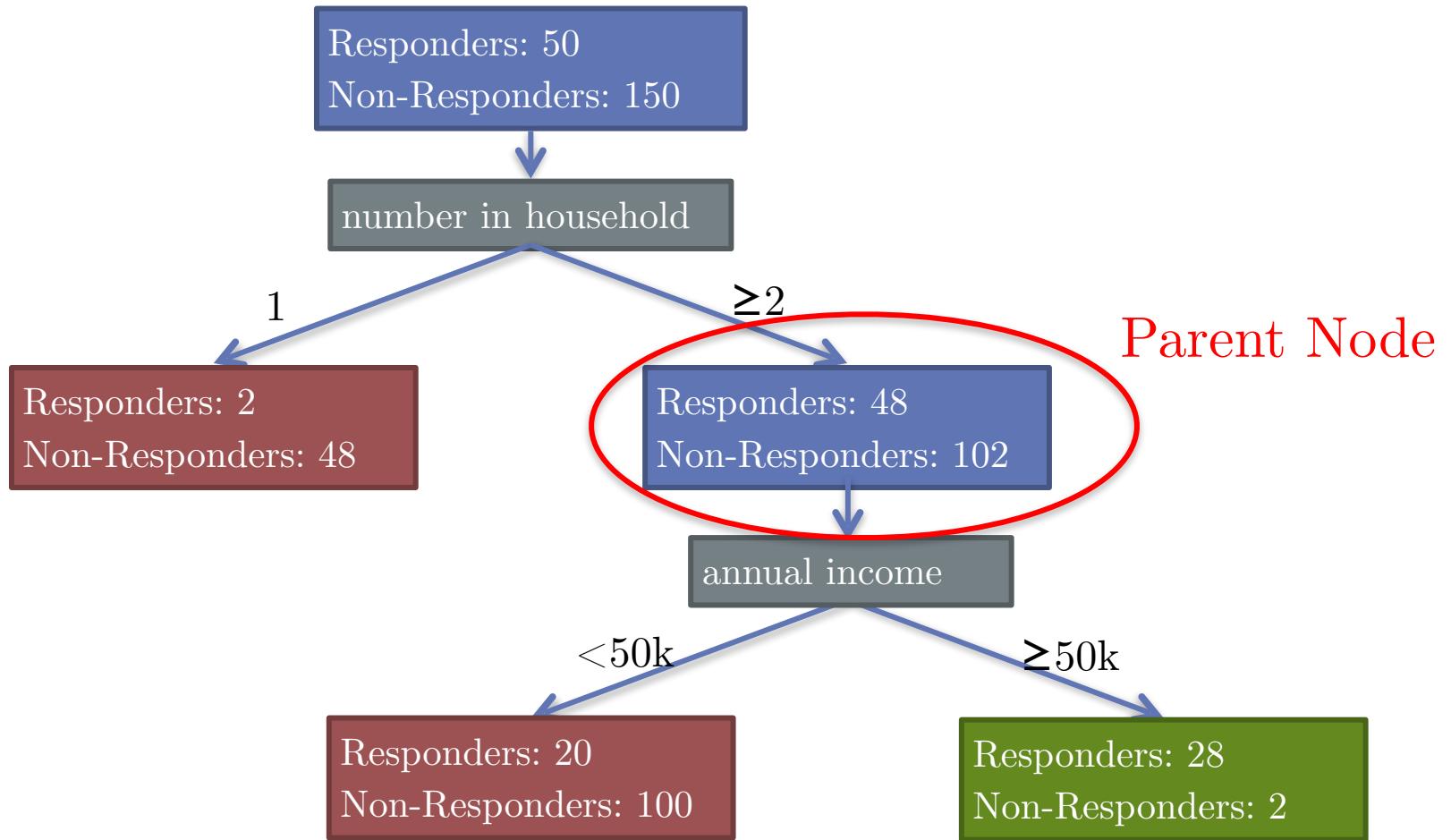
Decision Tree Model Creation



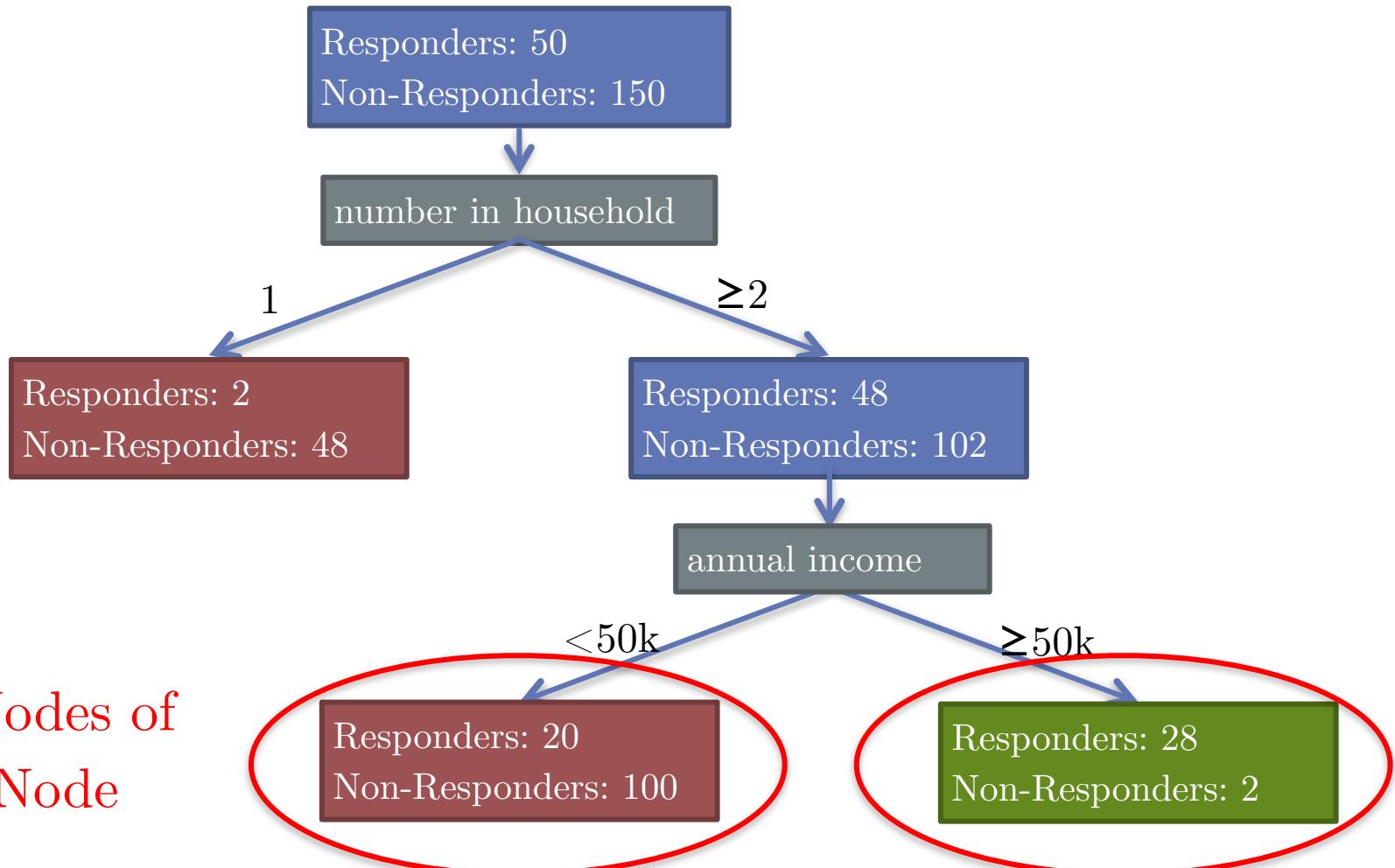
Decision Tree Model Creation



Decision Tree Model Creation



Decision Tree Model Creation



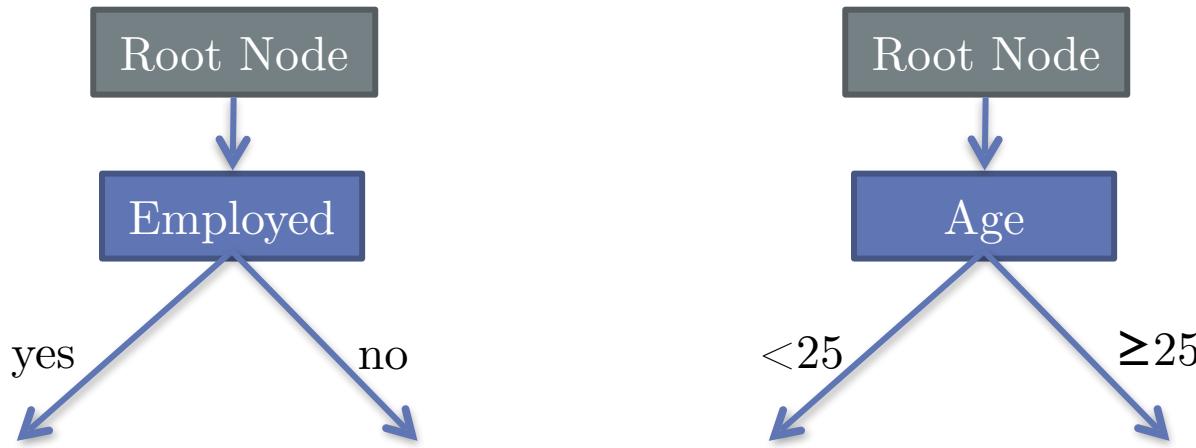
Classification Trees

• • •

Categorical/Ordinal Targets

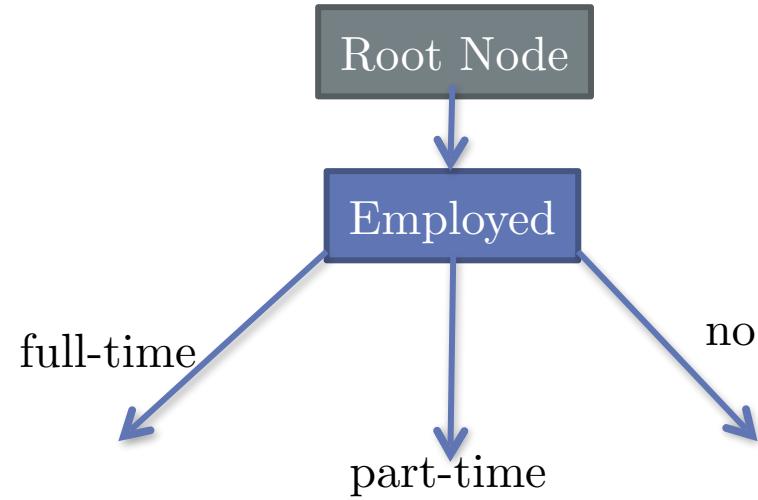
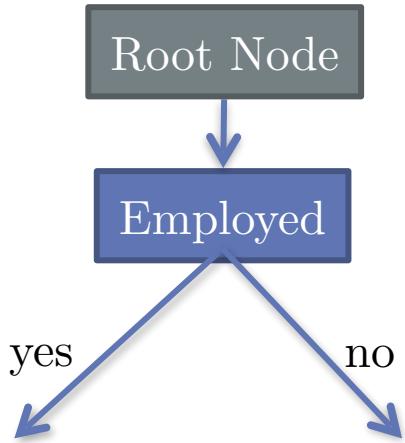
Building the model

- A tree is built by recursively partitioning the training data into successively **purer** subsets.
 - (Having mostly No's **or** mostly Yes's for the target.)
- Partitioning is done according to some condition.

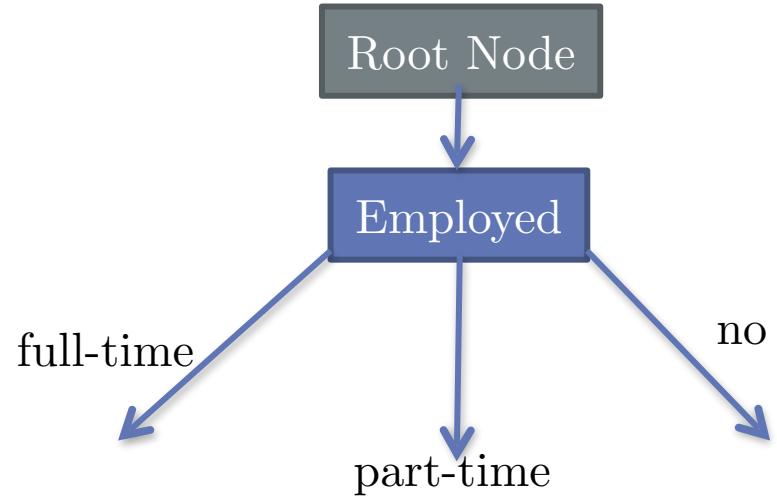
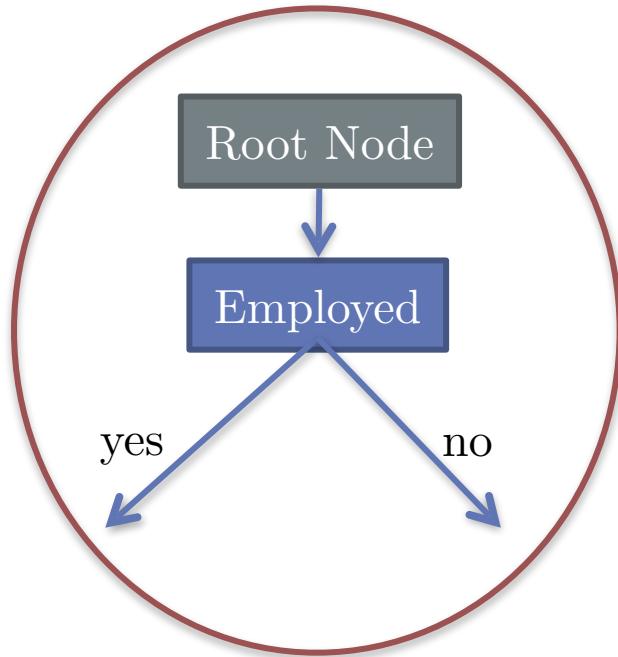


- How do we begin to assess these partitions?

Binary Splits vs. Multi-way Splits



Binary Splits vs. Multi-way Splits



- We will primarily discuss binary splits
- Everything is easily extended to multiway splits
- Binary trees are far more common

Categorical Input Variables

- We consider **every possible way** to separate into two distinct groups.
- Example:

Marital Status= {Single, Married, Other}

Leaf 1	Leaf 2
Single	Married, Other
Married	Single, Other
Other	Single, Married

- There are $2^{L-1} - 1$ possible splits for a variable with L levels

Ordinal Input Variables

- Only group together consecutive levels.
- Example:

Class = {Lower, Middle, Upper}

Leaf 1	Leaf 2
Lower	Middle, Upper
Lower, Middle	Upper

- There are L-1 such splits for an ordinal variable with L levels.

Continuous Input Variables

- Continuous Attributes: We consider all possible splits between data points *or bins of* the variable.
- Example:

Age={18,18,19,21,21,23,25,29,35,37,40,40,41,43}

Binary Splits

- Continuous Attributes: We consider all possible splits between data points or bins of the variable.
- Example:

Age = {18, 18, 19, 21, 21, 23, 25, 29, 35, 37, 40, 40, 41, 43}



Leaf 1	Leaf 2
Age < 19	Age \geq 19

Binary Splits

- Continuous Attributes: We consider all possible splits between data points *or bins of* the variable.
- Example:

Age = {18, 18, 19, 21, 21, 23, 25, 29, 35, 37, 40, 40, 41, 43}



Leaf 1	Leaf 2
Age < 21	Age \geq 21

Binary Splits

- Continuous Attributes: We consider all possible splits between data points *or bins of* the variable.
- Example:

Age = {18, 18, 19, 21, 21, 23, 25, 29, 35, 37, 40, 40, 41, 43}



Leaf 1	Leaf 2
Age < 23	Age \geq 23

Binary Splits

- Continuous Attributes: We consider all possible splits between data points *or bins of* the variable.
- Example:

Age = {18, 18, 19, 21, 21, 23, 25, 29, 35, 37, 40, 40, 41, 43}



Leaf 1	Leaf 2
Age < 25	Age ≥ 25

Binary Splits

- Continuous Attributes: We consider all possible splits between data points *or bins of* the variable.
- Example:

Age = {18, 18, 19, 21, 21, 23, 25, 29, 35, 37, 40, 40, 41, 43}

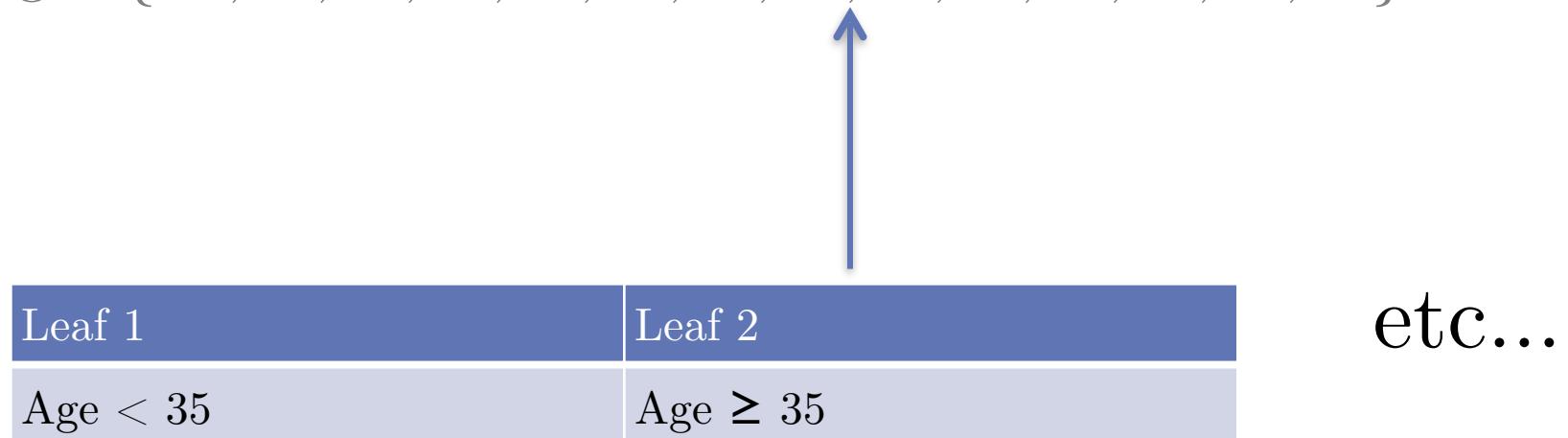


Leaf 1	Leaf 2
Age < 29	Age \geq 29

Binary Splits

- Continuous Attributes: We consider all possible splits between data points or bins of the variable.
- Example:

Age = {18, 18, 19, 21, 21, 23, 25, 29, 35, 37, 40, 40, 41, 43}

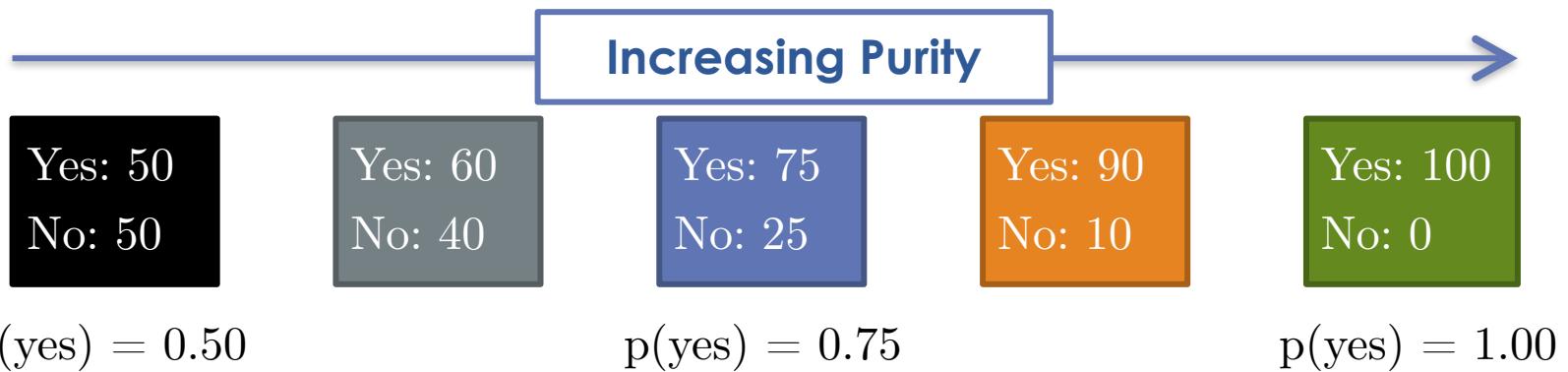


Missing Values

- One of the benefits of decision trees is their ability to handle missing values.
- Simply send missing values down one branch of the split (of course, it can get a lot fancier than that...)

Selecting the Best Split

- There are several measures used to select the best split.
- All are similar, but not identical
- All measure the **purity** of a node



- The more pure a leaf is, the less *training* error we make in that leaf.

Measures of Impurity

- Let $p(i | t) = p(class = i | node = t)$ be the fraction of records belonging to class i at a given node t . Let c be the number of classes in target variable.
- Entropy

$$Entropy(t) = - \sum_{i=1}^c p(i | t) \log_2 p(i | t)$$

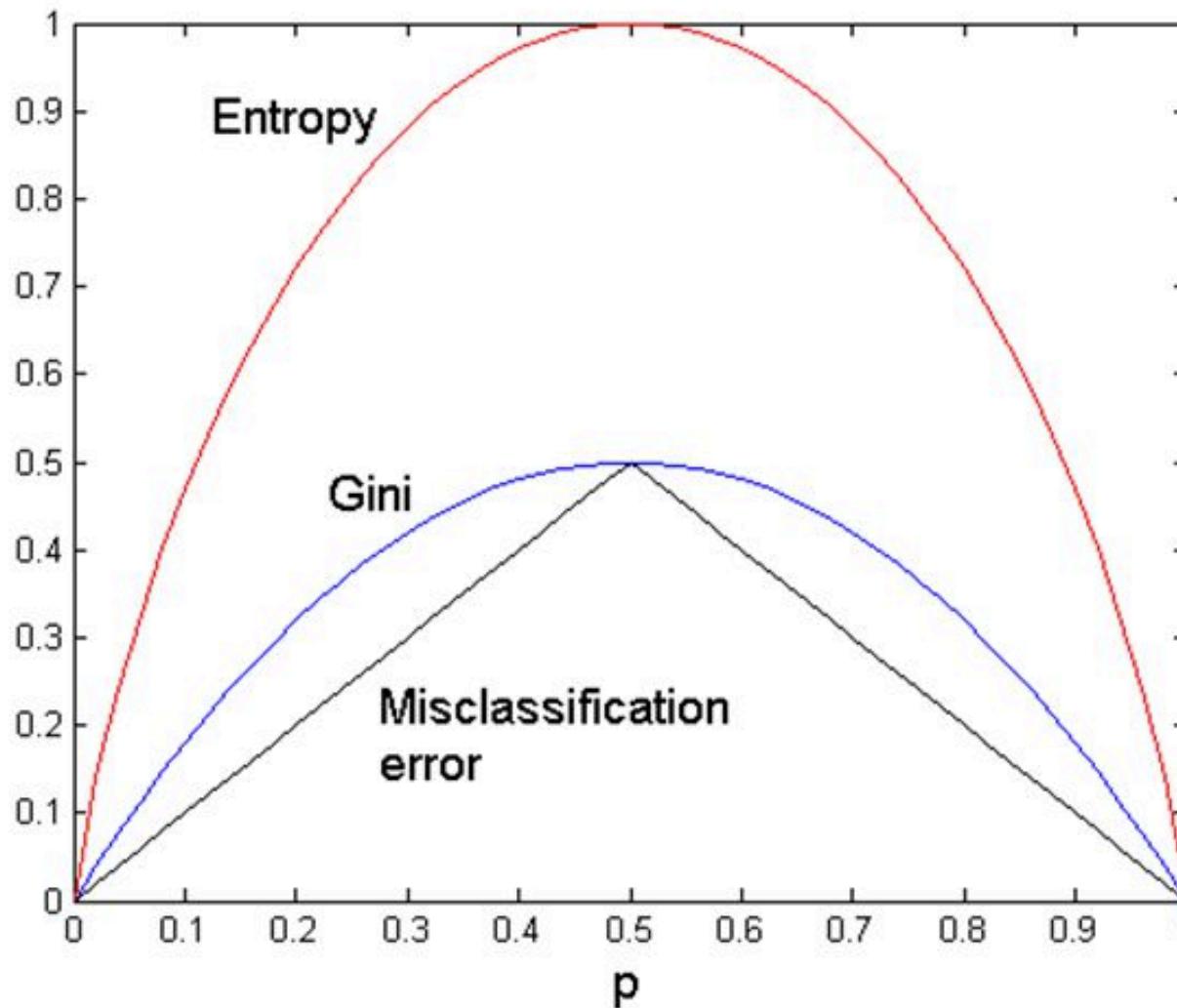
- Gini

$$Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

- Classification Error

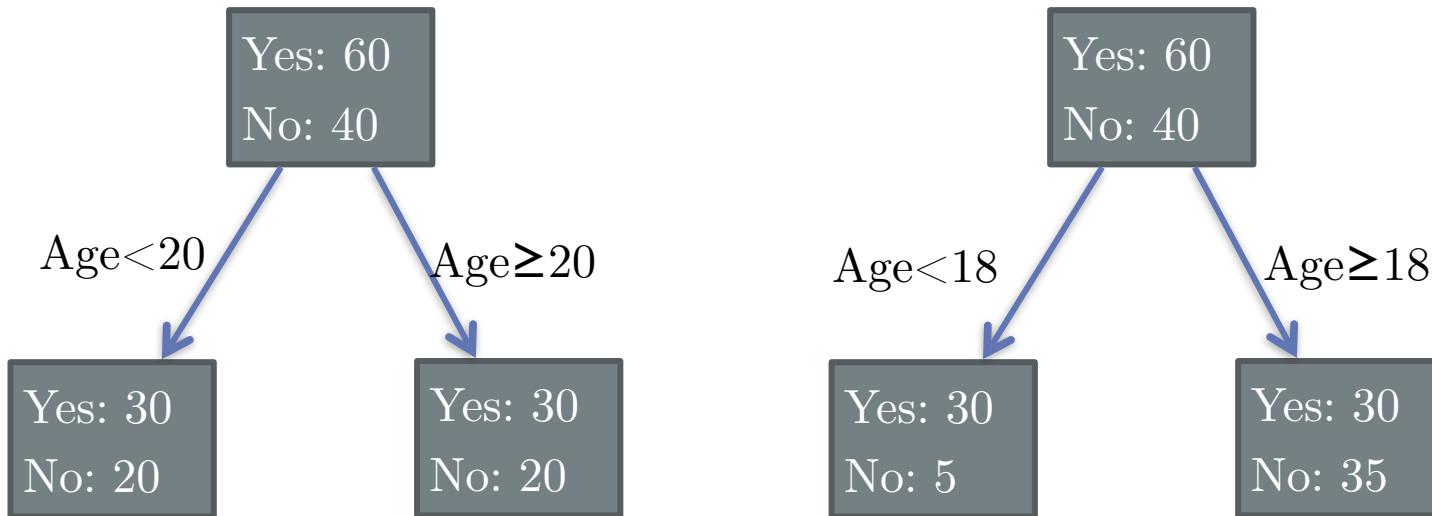
$$ClassificationError(t) = 1 - \max_i [p(i | t)]$$

Comparing Measures For a 2-class Problem



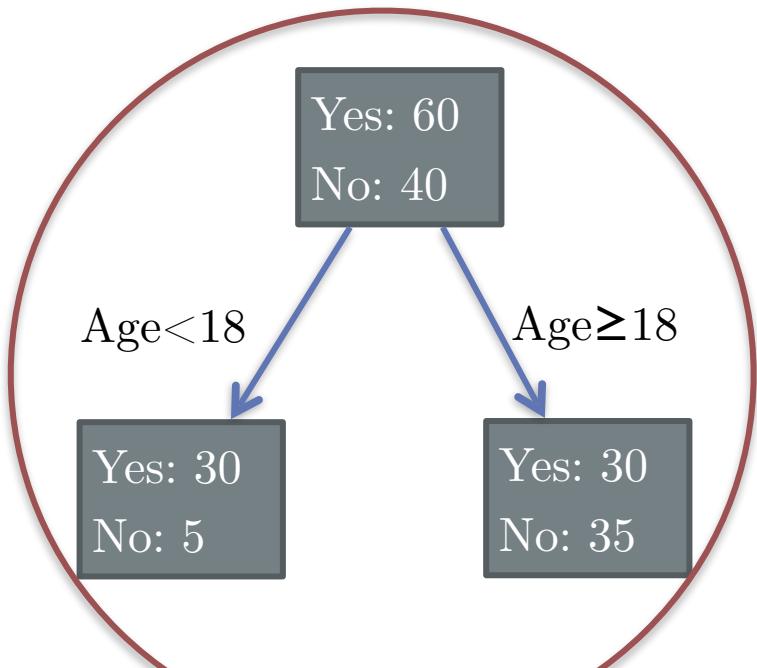
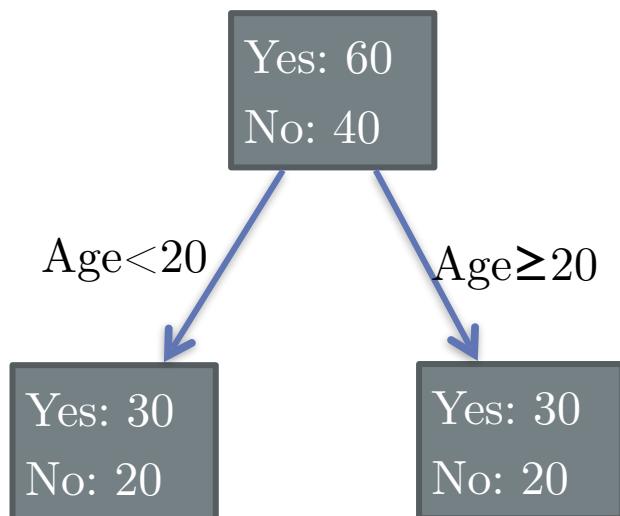
Selecting the best split

To assess a given test condition, we compare the impurity of the parent node (before split) with impurity of child nodes (after split).



Selecting the best split

To assess a given test condition, we compare the impurity of the parent node (before split) with impurity of child nodes (after split).



Split on the right has the best
GAIN in purity.
(i.e. Reduction of impurity)

Gain (Worth)

$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$

Δ := Gain

$I(t)$:= Impurity of parent node

$I(t_L)$ and $I(t_R)$:= Impurity of left/right child nodes

n := Number of observations in parent

n_L and n_R := Number of observations in left/right child

Gain (Worth)

$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$

Δ := Gain

$I(t)$:= Impurity of parent node

weighted avg. of
child node impurity

$I(t_L)$ and $I(t_R)$:= Impurity of left/right child nodes

n := Number of observations in parent

n_L and n_R := Number of observations in left/right child

Gain (Worth)

$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$

$\Delta :=$ Gain

$I(t) :=$ Impurity of parent node

$I(t_L)$ and $I(t_R) :=$ Impurity of left/right child nodes

$n :=$ Number of observations in parent

n_L and $n_R :=$ Number of observations in left/right child

Larger Gain → More pure branches

Gain (Worth)

$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$

When entropy is used, this difference in entropy is called *Information Gain*.

(For more information, see Tom Carter's slides at
<http://astarte.csustan.edu/~tom/SFI-CSSS/2005/info-lec.pdf>)

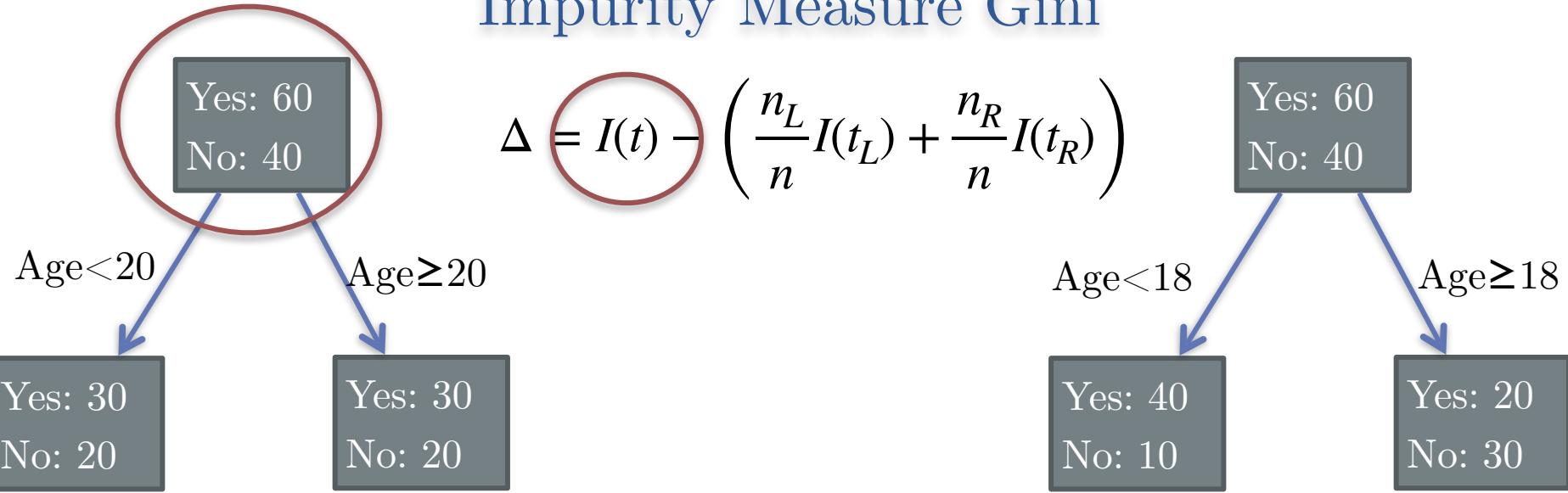
Example: Comparing 2 splits with Gain, Impurity Measure Gini



$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$

$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

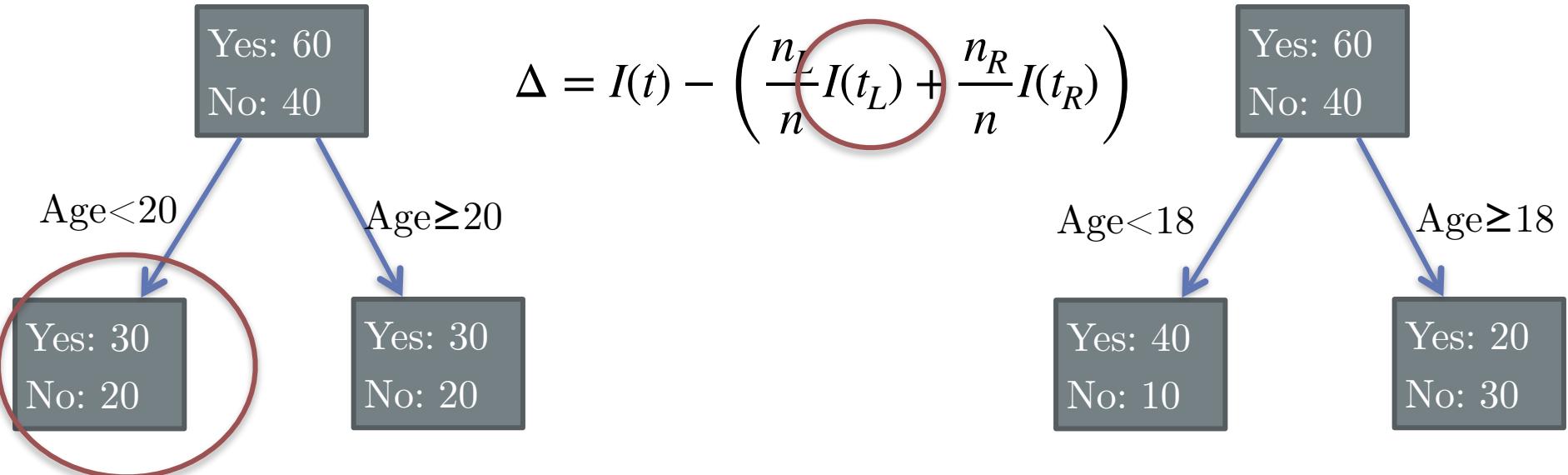
Example: Comparing 2 splits with Gain, Impurity Measure Gini



$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t) = 1 - \left[\left(\frac{60}{100} \right)^2 + \left(\frac{40}{100} \right)^2 \right] = 0.48$$

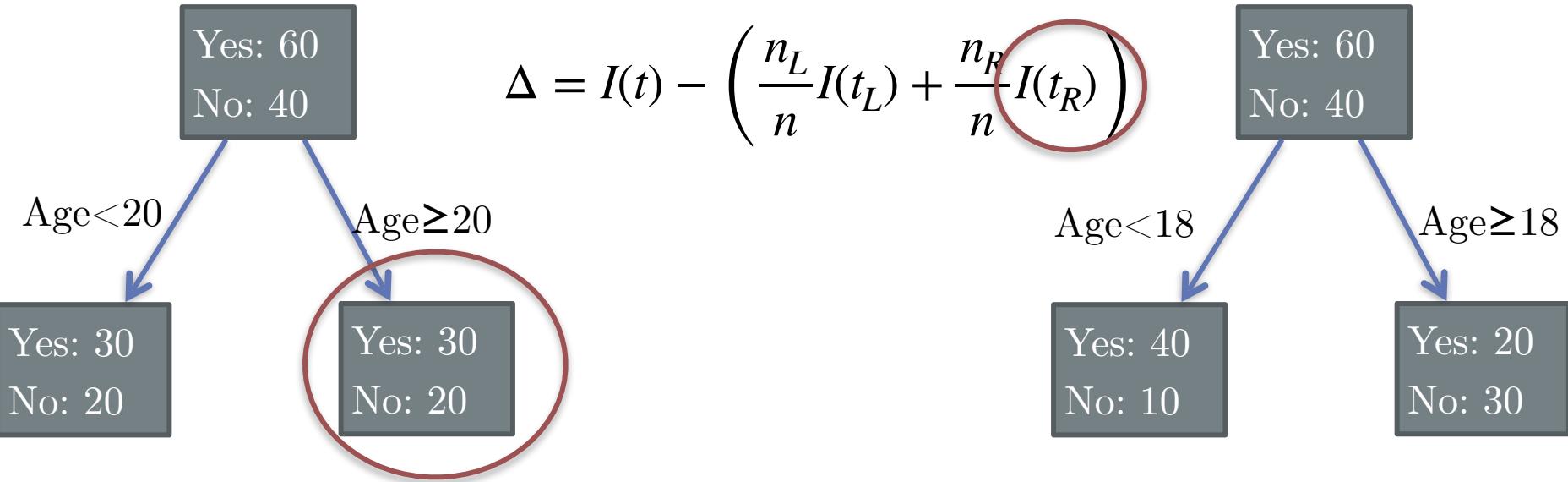
Example: Comparing 2 splits with Gain, Impurity Measure Gini



$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t_L) = 1 - \left[\left(\frac{30}{50} \right)^2 + \left(\frac{20}{50} \right)^2 \right] = 0.48$$

Example: Comparing 2 splits with Gain, Impurity Measure Gini

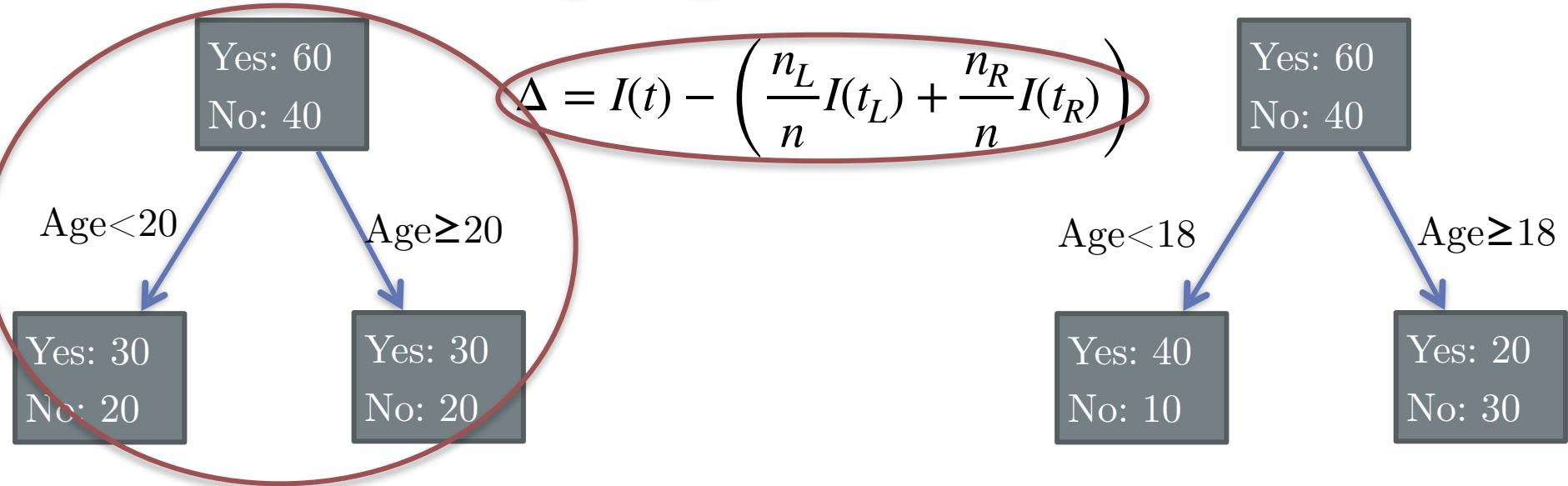


$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$

$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t_R) = 1 - \left[\left(\frac{30}{50} \right)^2 + \left(\frac{20}{50} \right)^2 \right] = 0.48$$

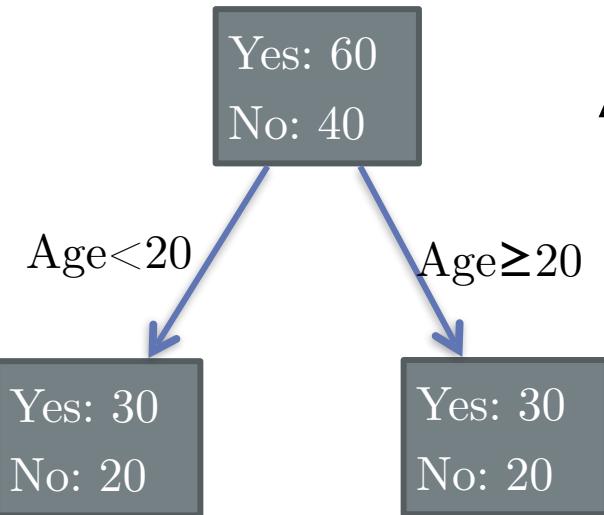
Example: Comparing 2 splits with Gain, Impurity Measure Gini



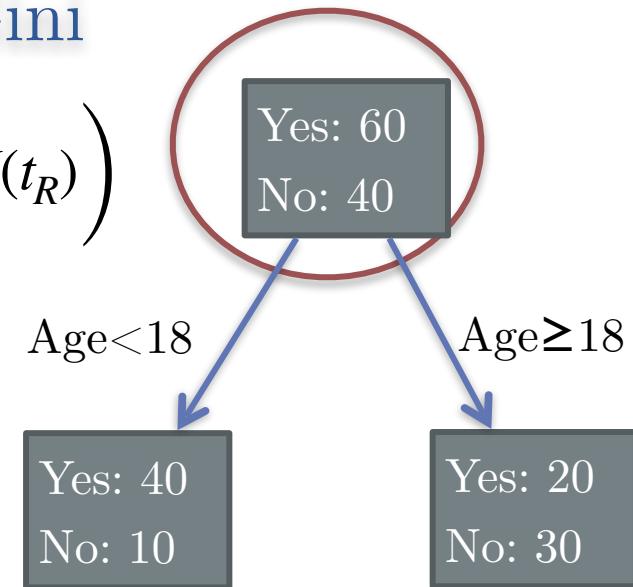
$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$\Delta = 0.48 - \left(\frac{50}{100} 0.48 + \frac{50}{100} 0.48 \right) = 0$$

Example: Comparing 2 splits with Gain, Impurity Measure Gini



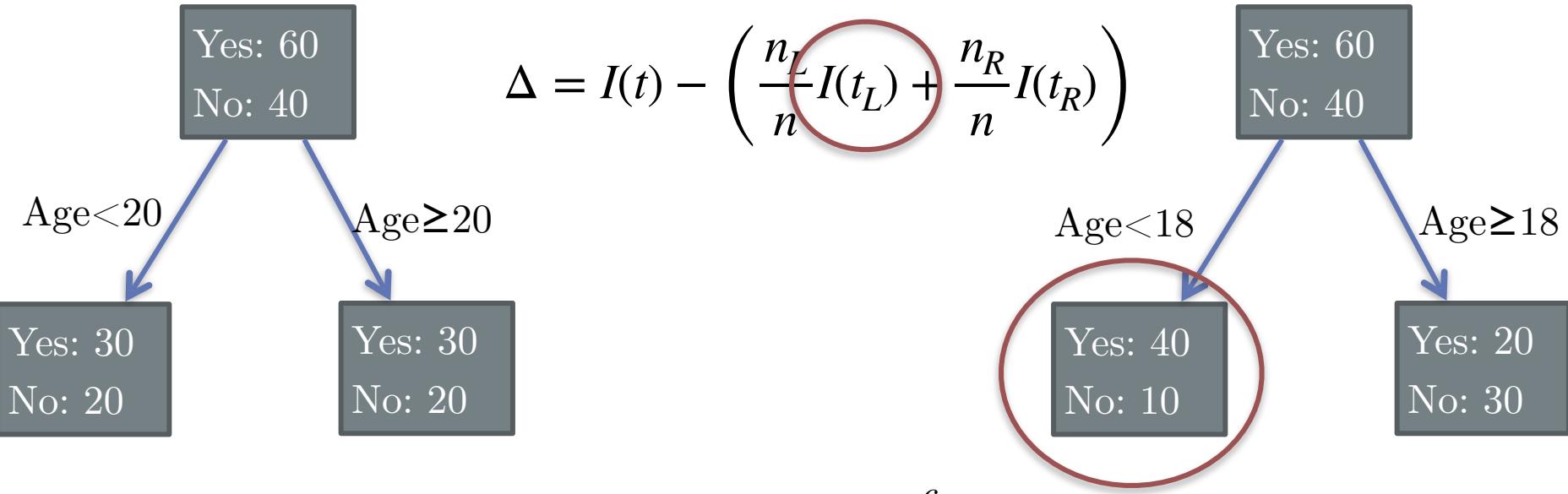
$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$



$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t) = 1 - \left[\left(\frac{60}{100} \right)^2 + \left(\frac{40}{100} \right)^2 \right] = 0.48$$

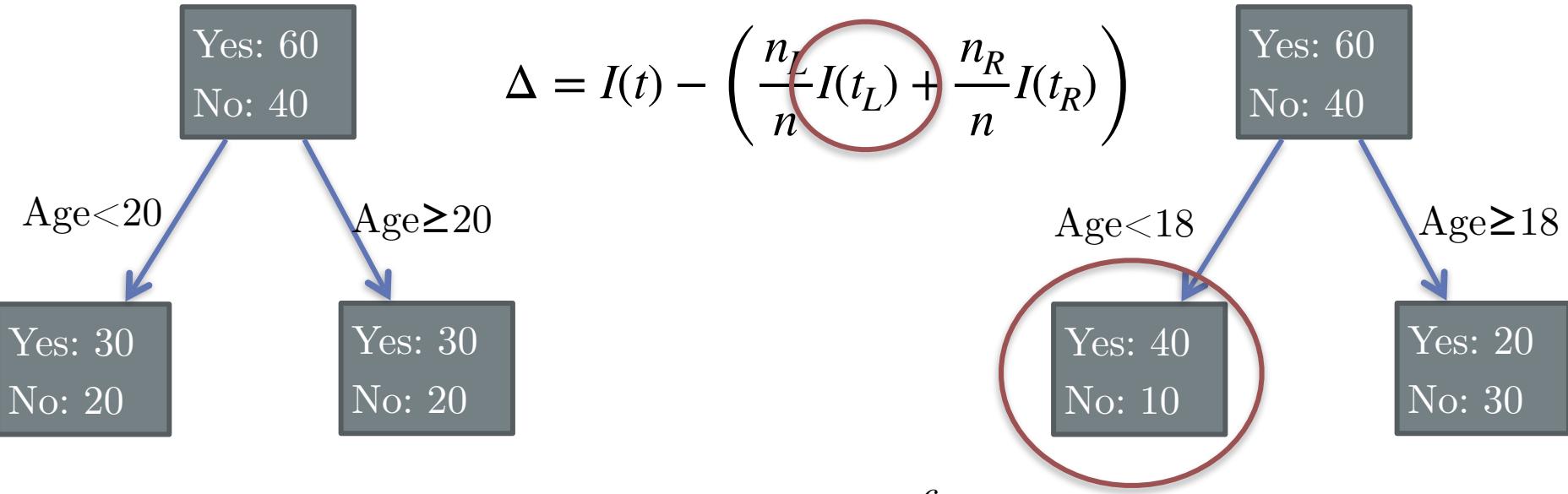
Example: Comparing 2 splits with Gain, Impurity Measure Gini



$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t_L) = 1 - \left[\left(\frac{40}{50} \right)^2 + \left(\frac{10}{50} \right)^2 \right] = 0.32$$

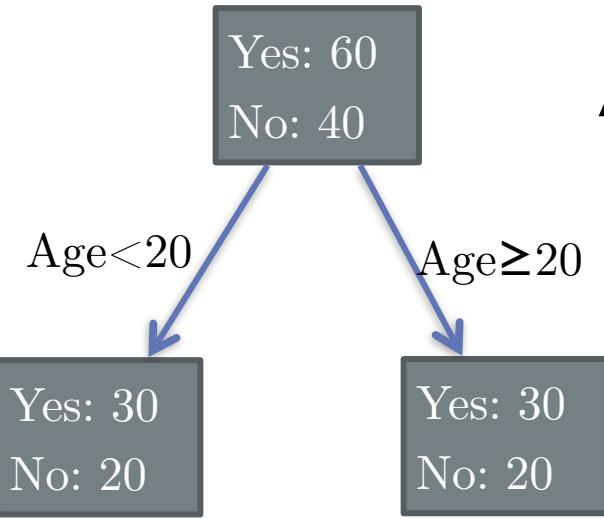
Example: Comparing 2 splits with Gain, Impurity Measure Gini



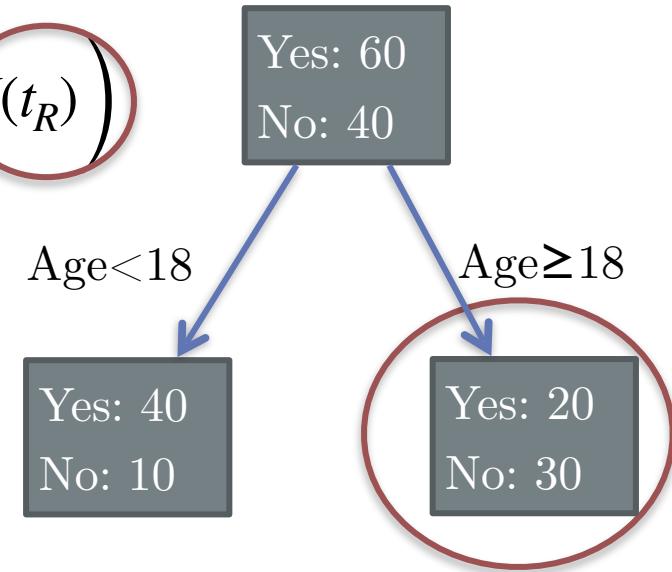
$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t_L) = 1 - \left[\left(\frac{40}{50} \right)^2 + \left(\frac{10}{50} \right)^2 \right] = 0.32$$

Example: Comparing 2 splits with Gain, Impurity Measure Gini



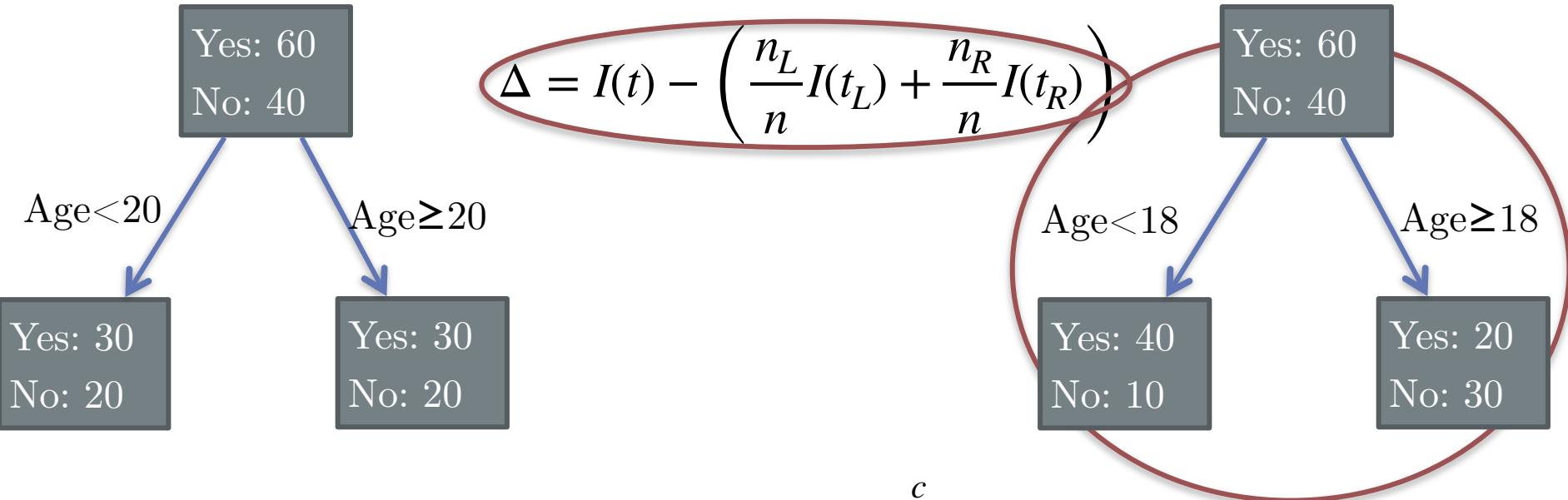
$$\Delta = I(t) - \left(\frac{n_L}{n} I(t_L) + \frac{n_R}{n} I(t_R) \right)$$



$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$I(t_R) = 1 - \left[\left(\frac{20}{50} \right)^2 + \left(\frac{30}{50} \right)^2 \right] = 0.48$$

Example: Comparing 2 splits with Gain, Impurity Measure Gini



$$I(t) = Gini(t) = 1 - \sum_{i=1}^c [p(i | t)]^2$$

$$\Delta = 0.48 - \left(\frac{50}{100} 0.32 + \frac{50}{100} 0.48 \right) = 0.08$$

So the split on the right has a higher gain and
is thus the better split

Creating the tree

- Compute the gain for all possible splits and select the best one.
- Repeat process recursively until some stopping condition is met
 - No splits meet some minimum Gain
 - All leaves have some minimum number of observations
 - A stopping condition is a way of *prepruning* the tree
- Prune Tree
 - Generally difficult to choose the right thresholds in prepruning
 - Can grow a larger tree and prune back branches in supervised fashion. (Essentially picking the threshold after the fact.)

Pruning a Decision Tree

- Simplifies the model
 - Occam's razor – law of parsimony
 - "Plurality is not to be posited without necessity"
(Duns Scotus 1290)
- Prevents overfitting the training data
 - An accurate model on training: one bin for each leaf!
#TerribleIdea
- **Simply remove leaves/nodes** in a bottom-up fashion, cutting splits with lowest gain first, while **optimizing performance on validation data**

Viya Demo 1

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Telco Customer Churn

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<https://www.kaggle.com/blastchar/telco-customer-churn>

Problem Introduction

Goal: Predict behavior to retain customers. Analyze all relevant customer data and develop focused customer retention programs.

The data set includes information about:

- Customers who left within the last month (and customers who did not)
 - the **target column** is called **Churn**
- **Services that each customer has signed up for** – *phone, multiple lines, internet, online security, online backup, device protection, tech support, and streaming TV and movies*
- Customer **account information** – *tenure as a customer, contract, payment method, paperless billing, monthly charges, and total charges*
- **Demographic info** about customers – *gender, age range, and if they have partners and dependents*

1

ANALYTICS LIFE CYCLE

- Manage Data
- Prepare Data
- Explore and Visualize**
- Build Models
- Manage Models
- Share and Collaborate
- Develop SAS Code

Objects 2

Filter

Standard container

Content

- Data-driven content
- Image
- Text
- Web content

SAS Visual Statistics

- Cluster
- Decision tree
- Generalized additive model
- Generalized linear model
- Linear regression
- Logistic regression
- Model comparison
- Nonparametric logistic regression

SAS Visual Data Mining and Machine L...

... 3

Data

TELCOCHURN

Filter

+ New data item

Sug...

Out...

Hierarchy

Custom category

Calculated item

Geography item

Parameter

Interaction effect

Spline effect

Partition

New Partition

Name: Partition

Based on:

Data item Sampling

Sampling method:

Simple random sampling

Number of partitions:

2

Training partition sampling percentage: * 80

Random number seed

Random seed: * 11117

OK Cancel

4

Data Roles

Decision tree - Churn 2

Response

Churn

5

Predictors

- Contract
 - Dependents
 - DeviceProtection
 - gender
 - InternetService
 - MultipleLines
 - OnlineBackup
 - OnlineSecurity
 - PaperlessBilling
 - Partner
 - PaymentMethod
 - PhoneService
 - StreamingMov...
 - StreamingTV
 - TechSupport
 - MonthlyCharges
 - SeniorCitizen
 - tenure
 - TotalCharges
- + Add
- Partition ID
- Training

Options

Roles

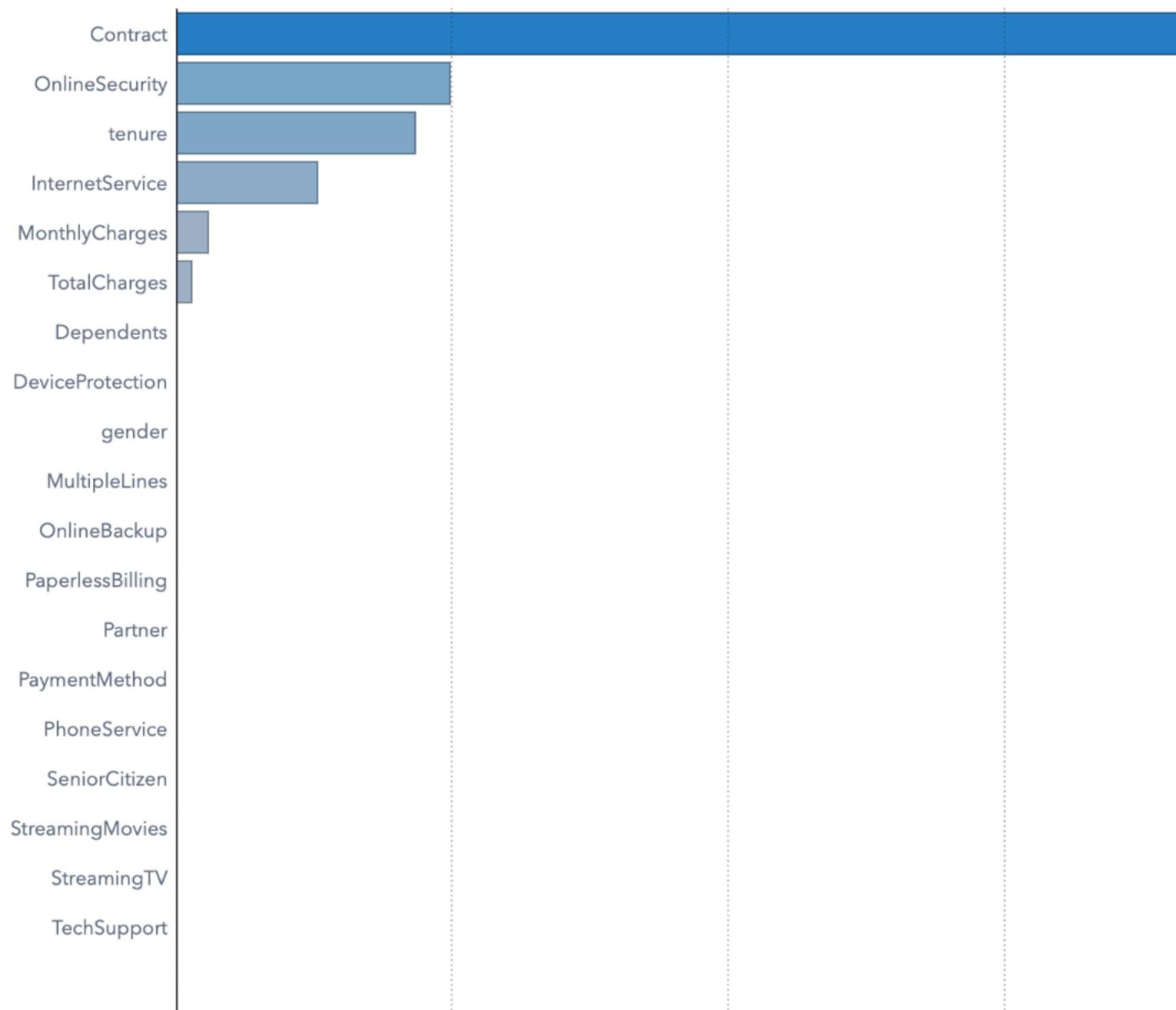
Actions

Rules

Filters

Ranks

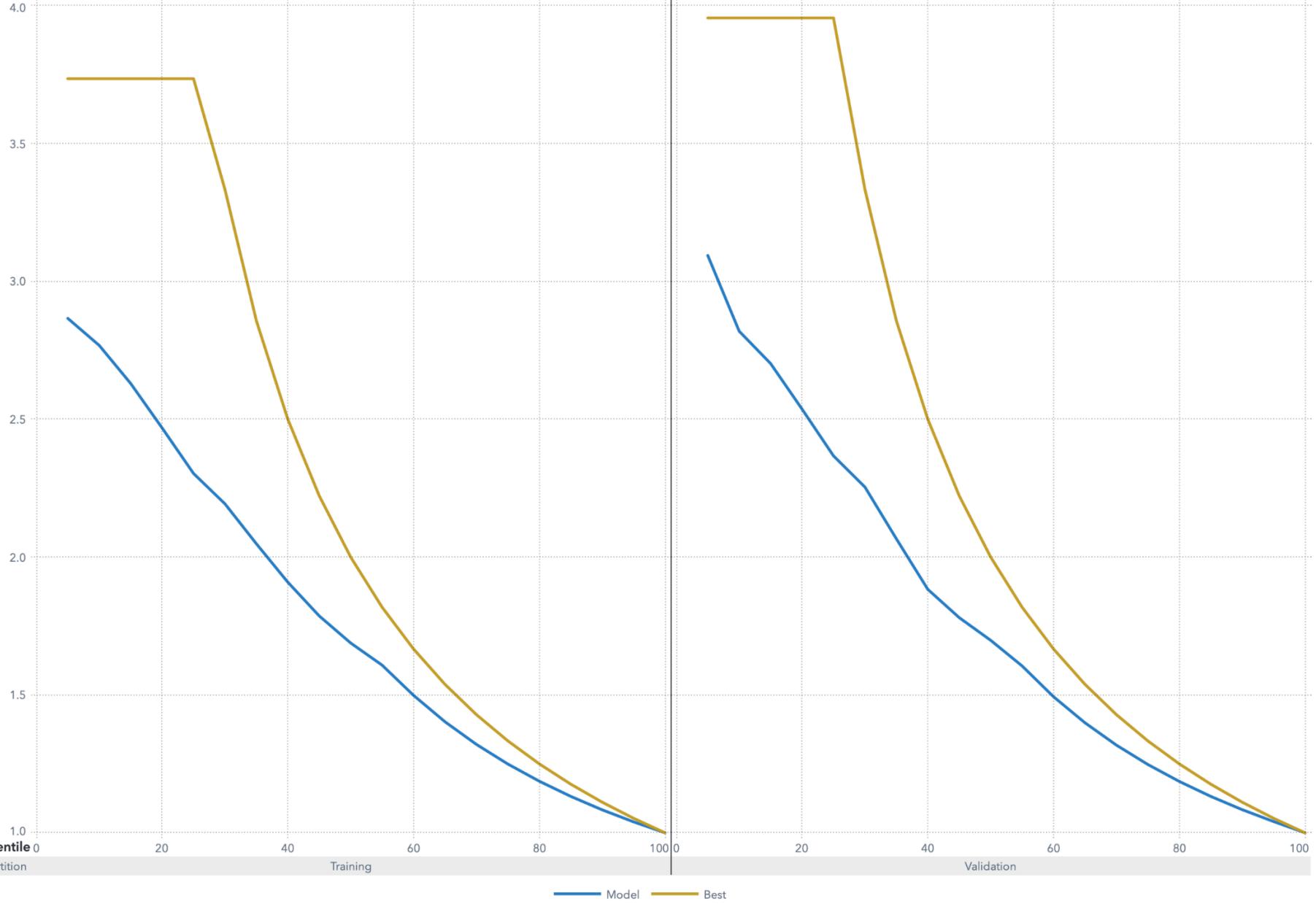
Variable Importance

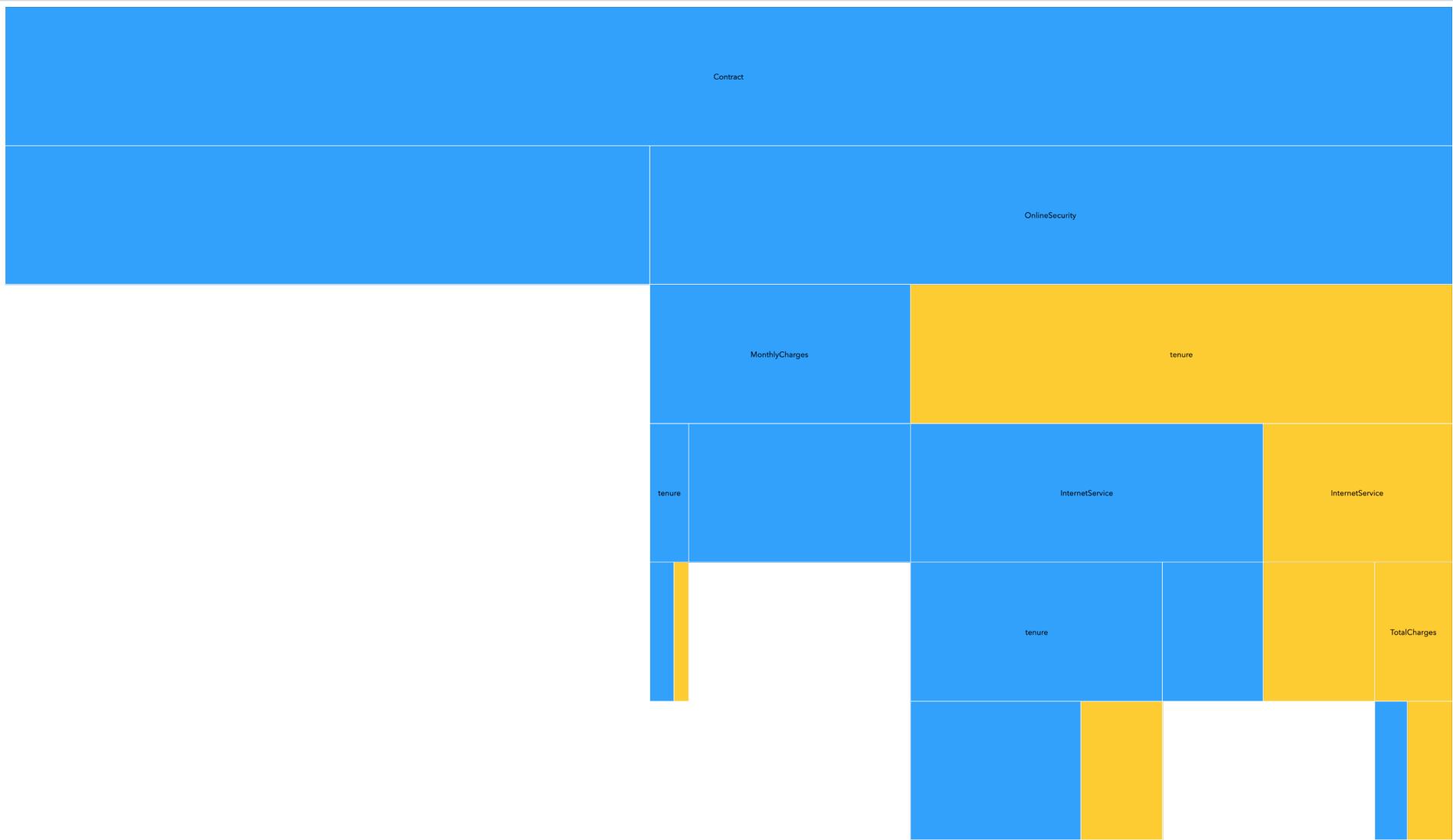


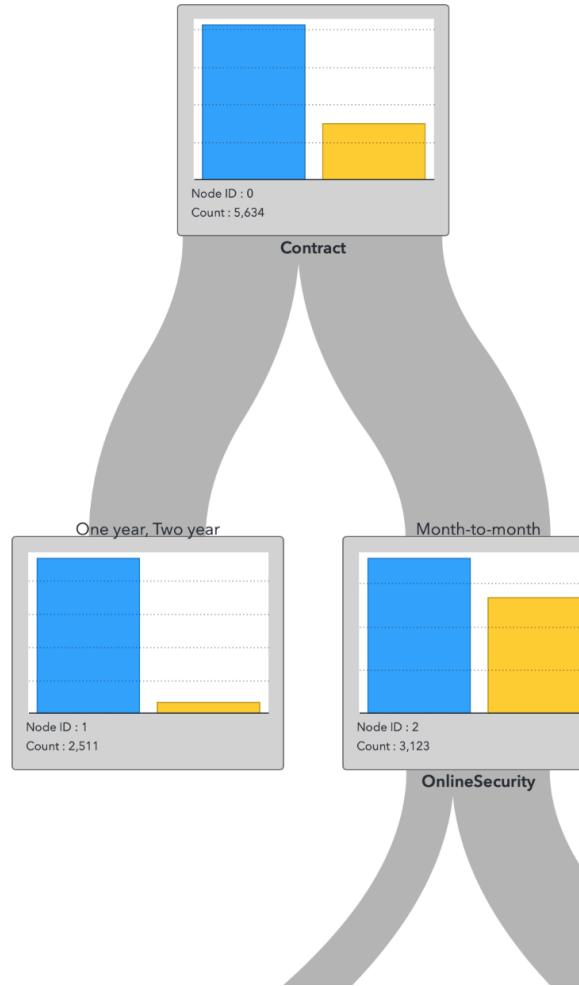
Lift

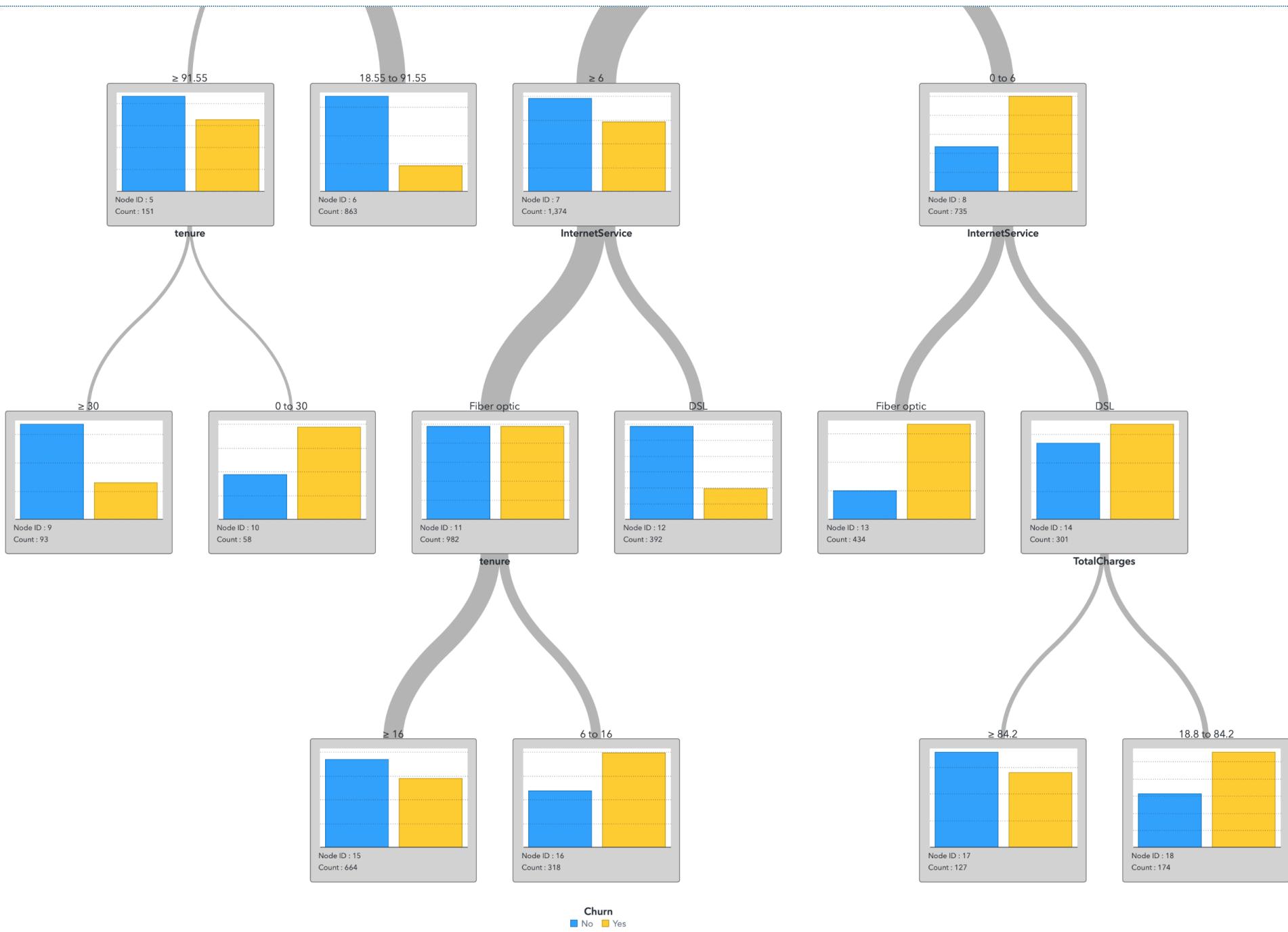


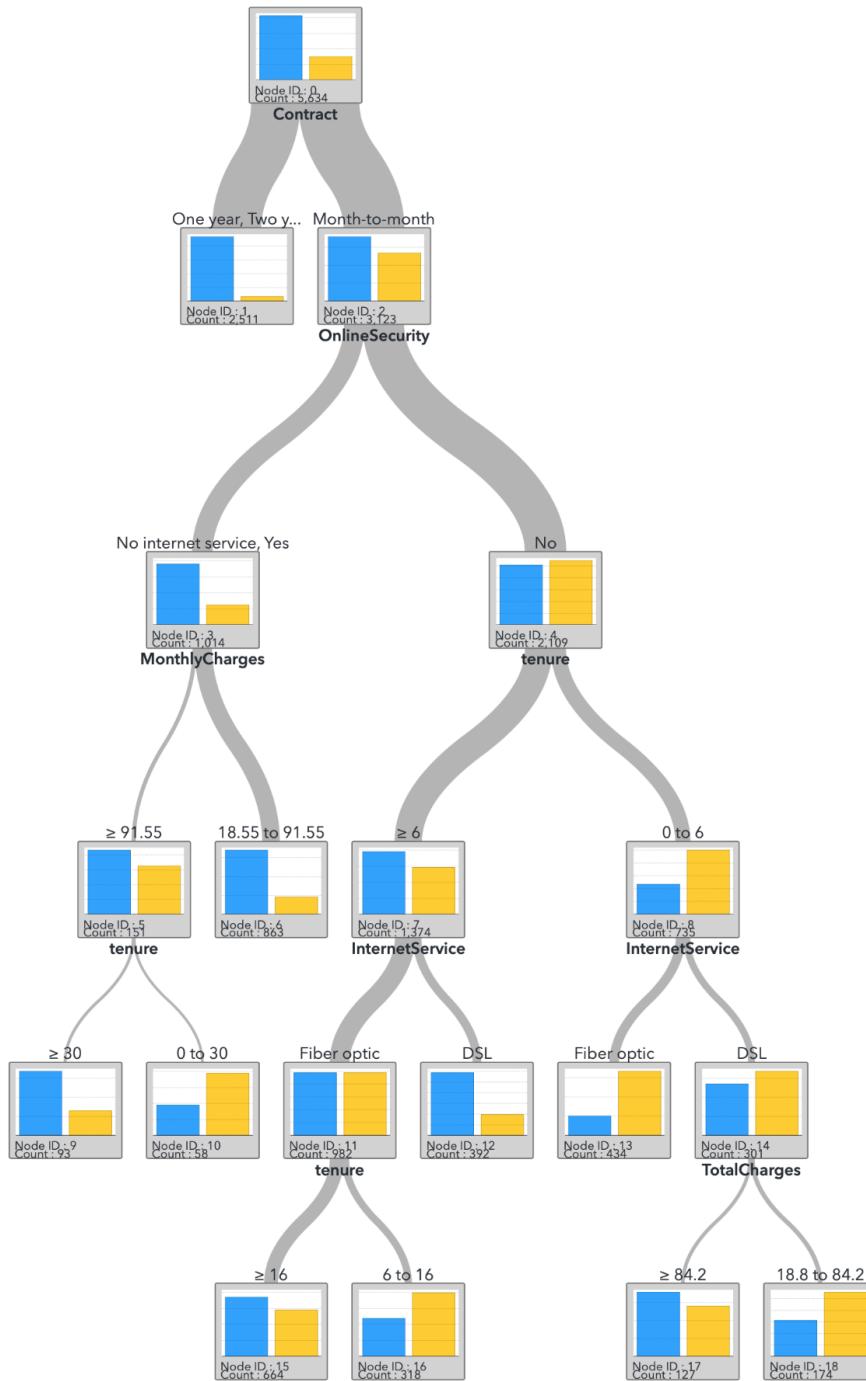
Cumulative Lift











Part II

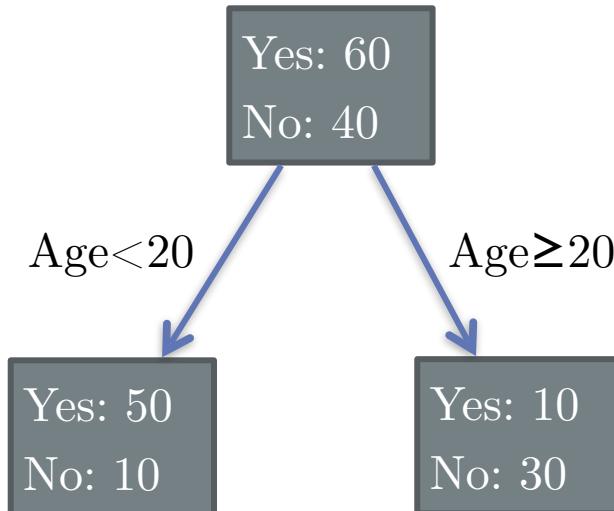
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CHAID and Regression Trees

CHAID

CHi-squared Automatic Interaction Detection

- 1980 PhD thesis by Gordon Kass
- Rather than using gain to determine splits, use chi-square tests!
- Analyze decision tree splits like we do contingency tables:

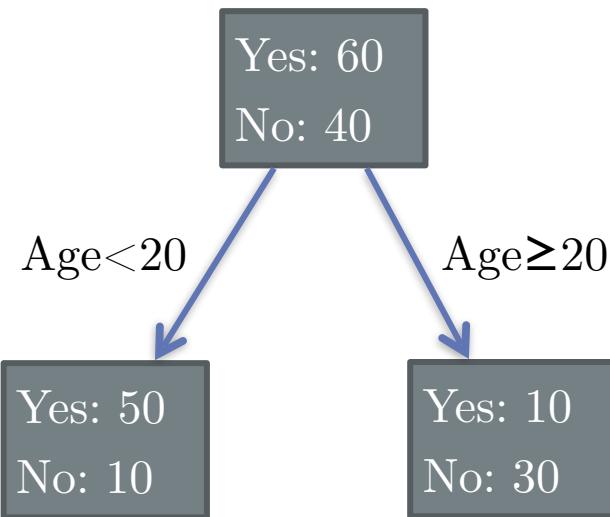


	Yes	No	Total
Age < 20	50	10	60
Age ≥ 20	10	30	40
Total	60	40	100

$$\chi^2 = \sum_{cells} \frac{(observed - expected)^2}{expected}$$

CHAID

CHi-squared Automatic Interaction Detection



	Yes	No	Total
Age < 20	50	10	60
Age ≥ 20	10	30	40
Total	60	40	100

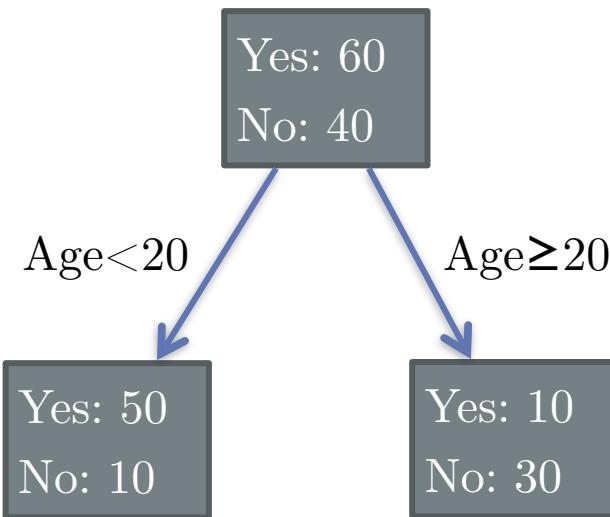
$$\chi^2 = \sum_{cells} \frac{(observed - expected)^2}{expected}$$

Larger χ^2 statistic \rightarrow Smaller p-value \rightarrow Stronger relationship

only b/c sample size is constant in comparison at a given parent node!

CHAID

CHi-squared Automatic Interaction Detection



	Yes	No	Total
Age < 20	50	10	60
Age ≥ 20	10	30	40
Total	60	40	100

$$\chi^2 = \sum_{cells} \frac{(observed - expected)^2}{expected}$$

Larger χ^2 statistic \rightarrow Smaller p-value \rightarrow Stronger relationship

Uses **logworth** to choose a split: $\text{logworth}(p) = -\log_{10}(p)$

Logworth

$$\text{logworth}(p) = -\log_{10}(p)$$

Tells us approx # of decimal places of our p-value.

Examples:

- $\text{logworth}(0.001) = -\log_{10}(0.001) = -(-3) = 3$.
- $\text{logworth}(0.0001) = 4$
- $\text{logworth}(0.0004)$ is between 3 and 4
 - $0.0001 < 0.0004 < 0.001$
 - $\log_{10}(0.0001) < \log_{10}(0.0004) < \log_{10}(0.001)$
 - $-\log_{10}(0.0001) > -\log_{10}(0.0004) > -\log_{10}(0.001)$
 - $4 > -\log_{10}(0.0004) > 3$

LARGER LOGWORTH => BETTER SPLIT.

Kass Adjustments (i.e. Bonferroni Adjustments)

- Hypothesis testing to compare many variables at many potential splits. (Could be thousands of comparisons!)
- Beware the family-wise error rate!!
- **Adjust the test significance to (α/m)** where α is your desired significance level and m is number of tests.
- **Equivalent to multiplying p-values by m and keeping α unchanged.**

Kass Adjustments (i.e. Bonferroni Adjustments)

Suppose we compare *Age* (interval) with *Insurance Status* (binary).

No Adjustment

- best p-value for *Age* is **0.01** and occurs when splitting at $\text{Age} < 20$, $\text{Age} \geq 20$
- p-value for *Insurance Status* is **0.05**

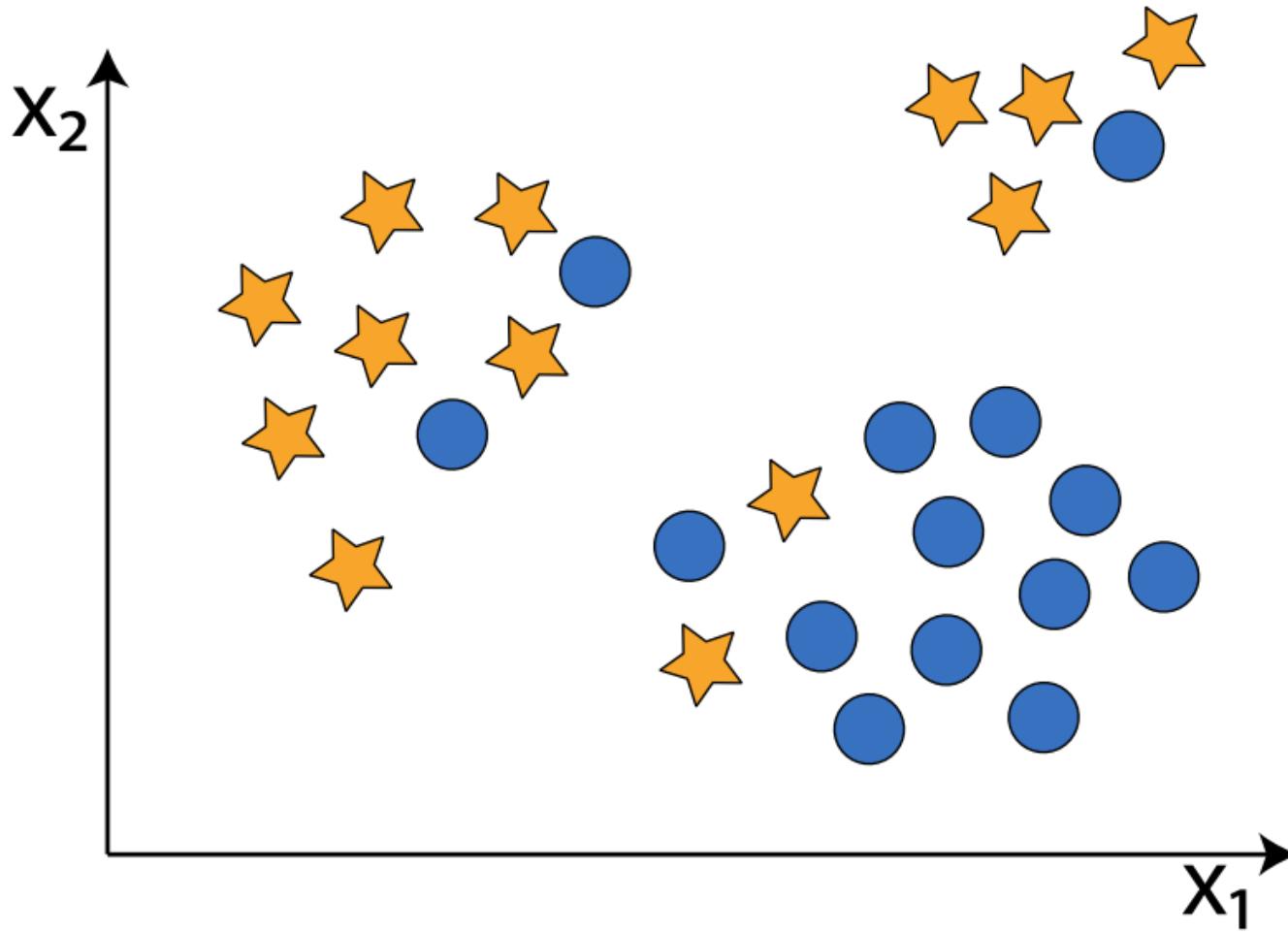
Pick
 $\text{Age} < 20$, $\text{Age} \geq 20$
as the splitting
criterion.

Bonferroni Adjustment

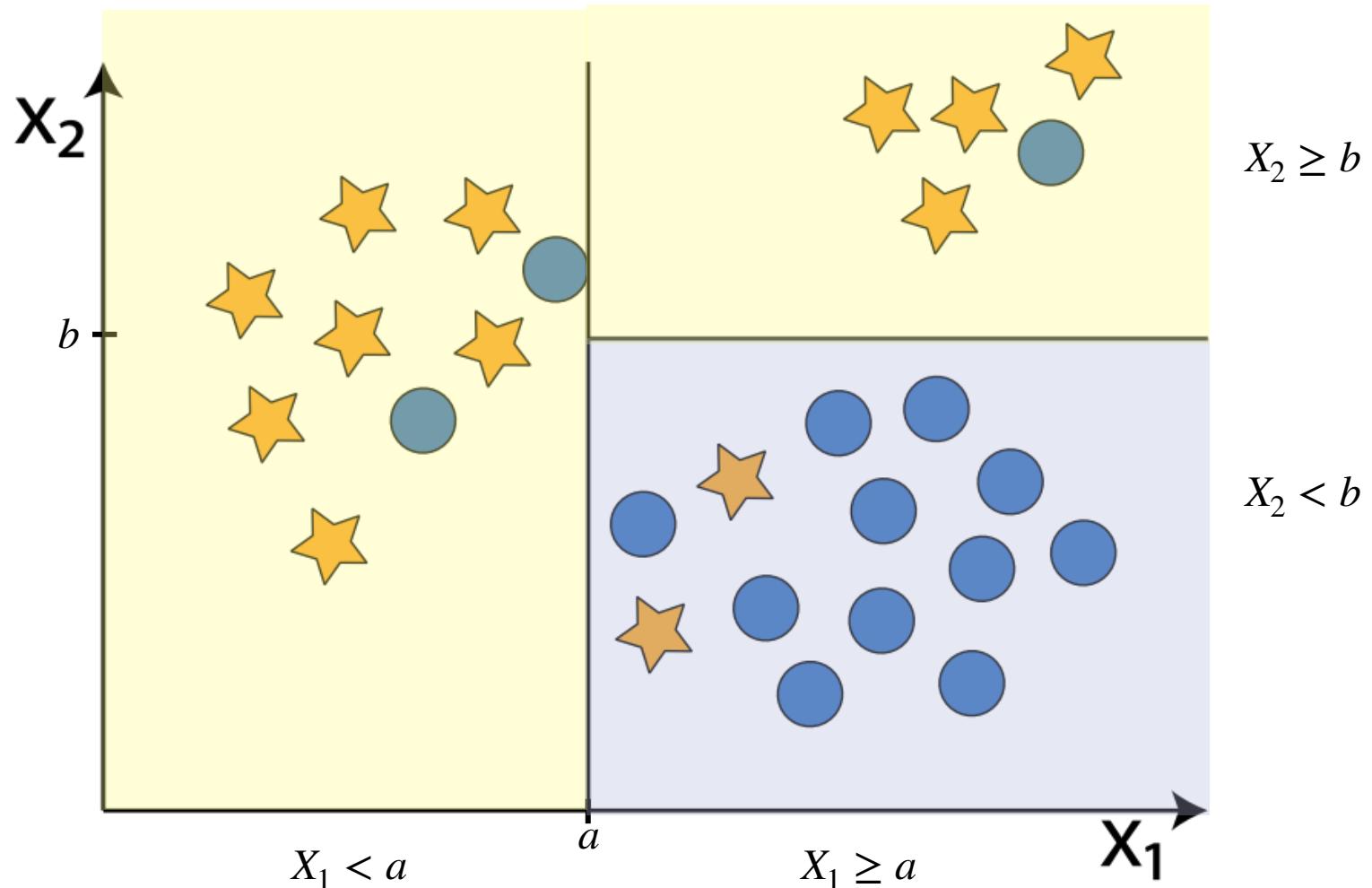
- Age had 51 unique values (50 possible splits)
- Insurance Status had 1
- Not fair to compare these p-values! In 50 tests, using **one** with a p-value of 0.01 is not convincing!
- Adjust p-values by multiplying by number of tests:
 - Age: $(0.01)^*50 = 0.5$
 - Insurance Status: $(0.05)^*1 = 0.05$

Pick
Insurance Status
as splitting criterion.

Decision Tree Boundaries

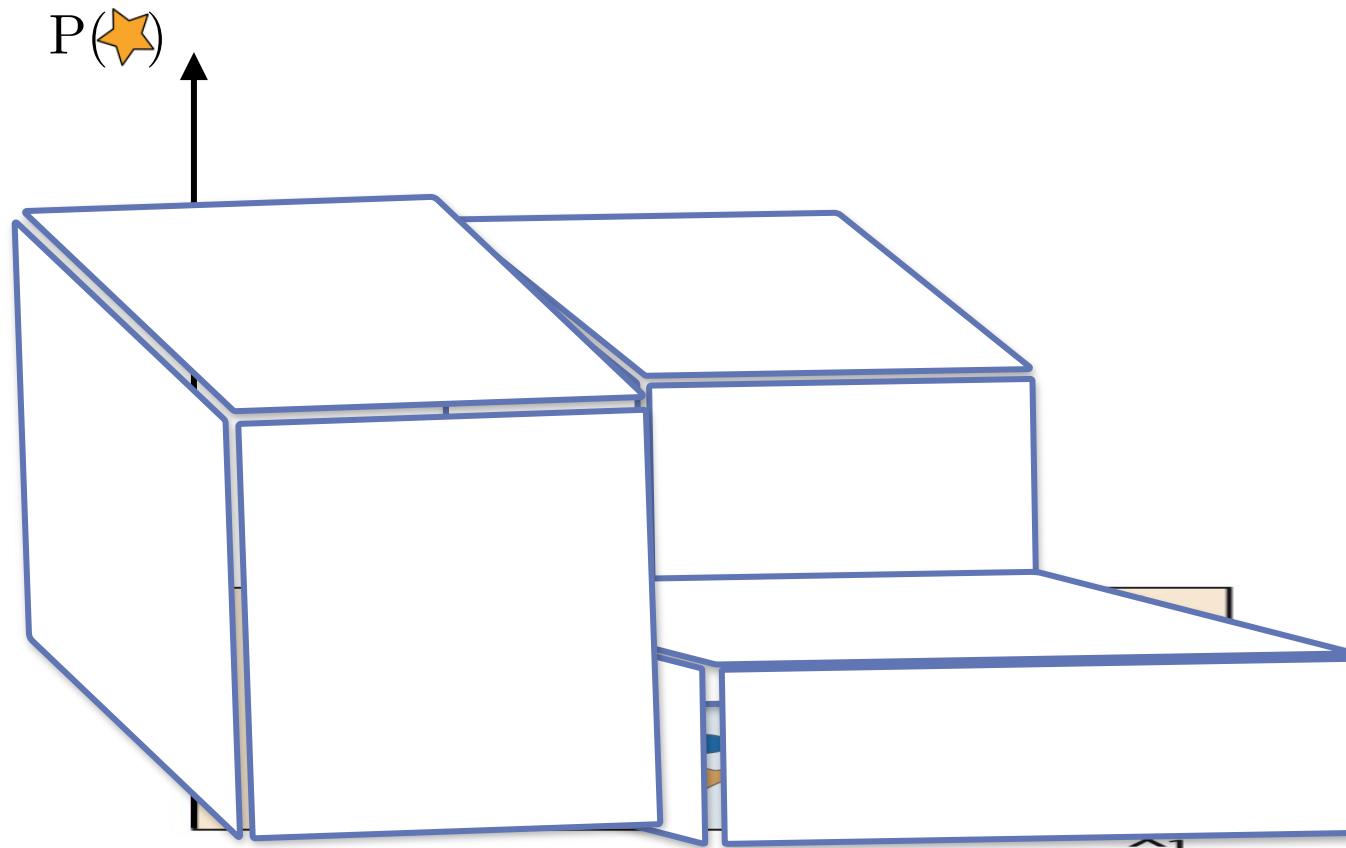


Decision Tree Boundaries



Decision Tree Response Surface

(Building with legos - no diagonals!)

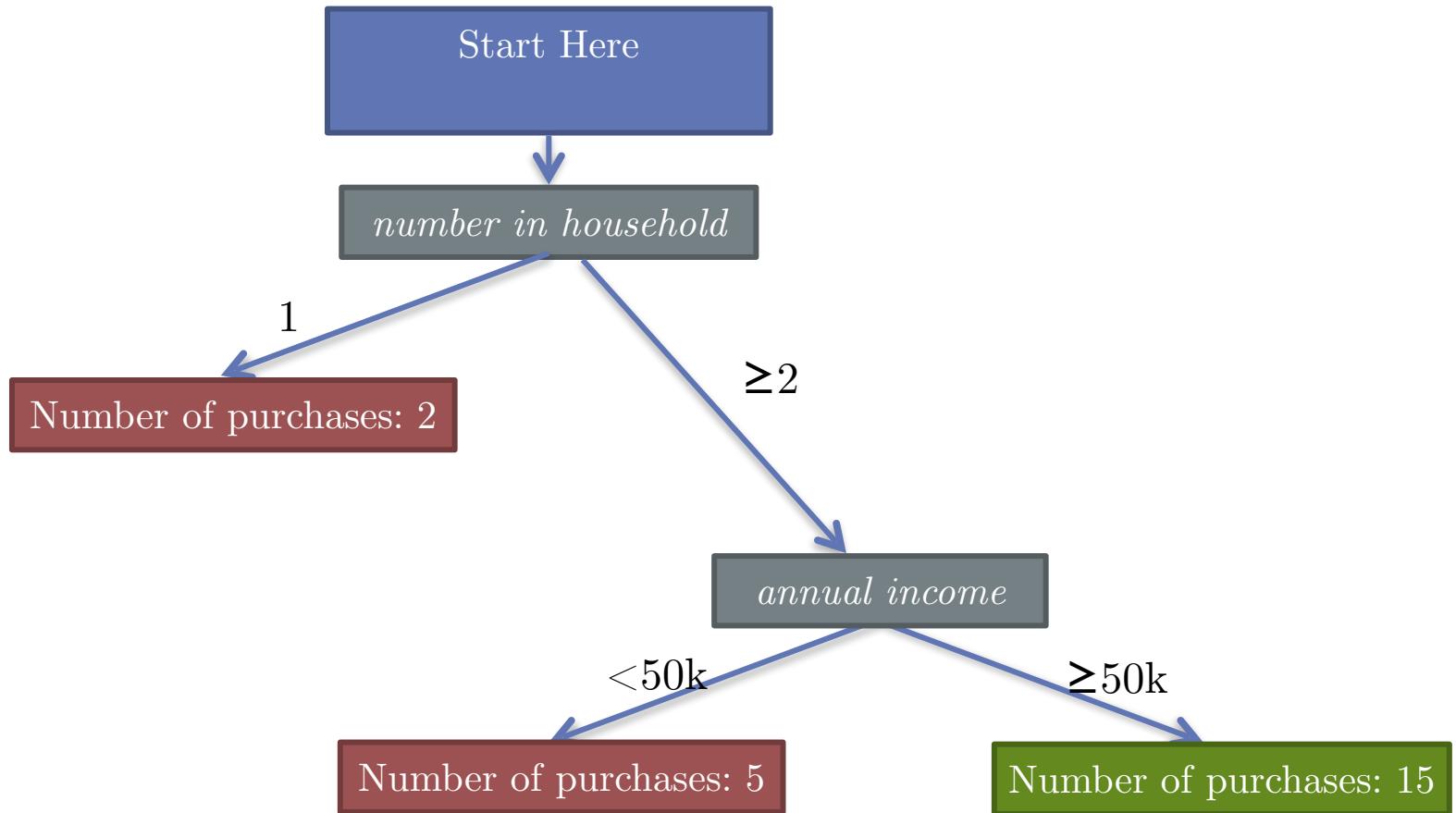


Regression Trees

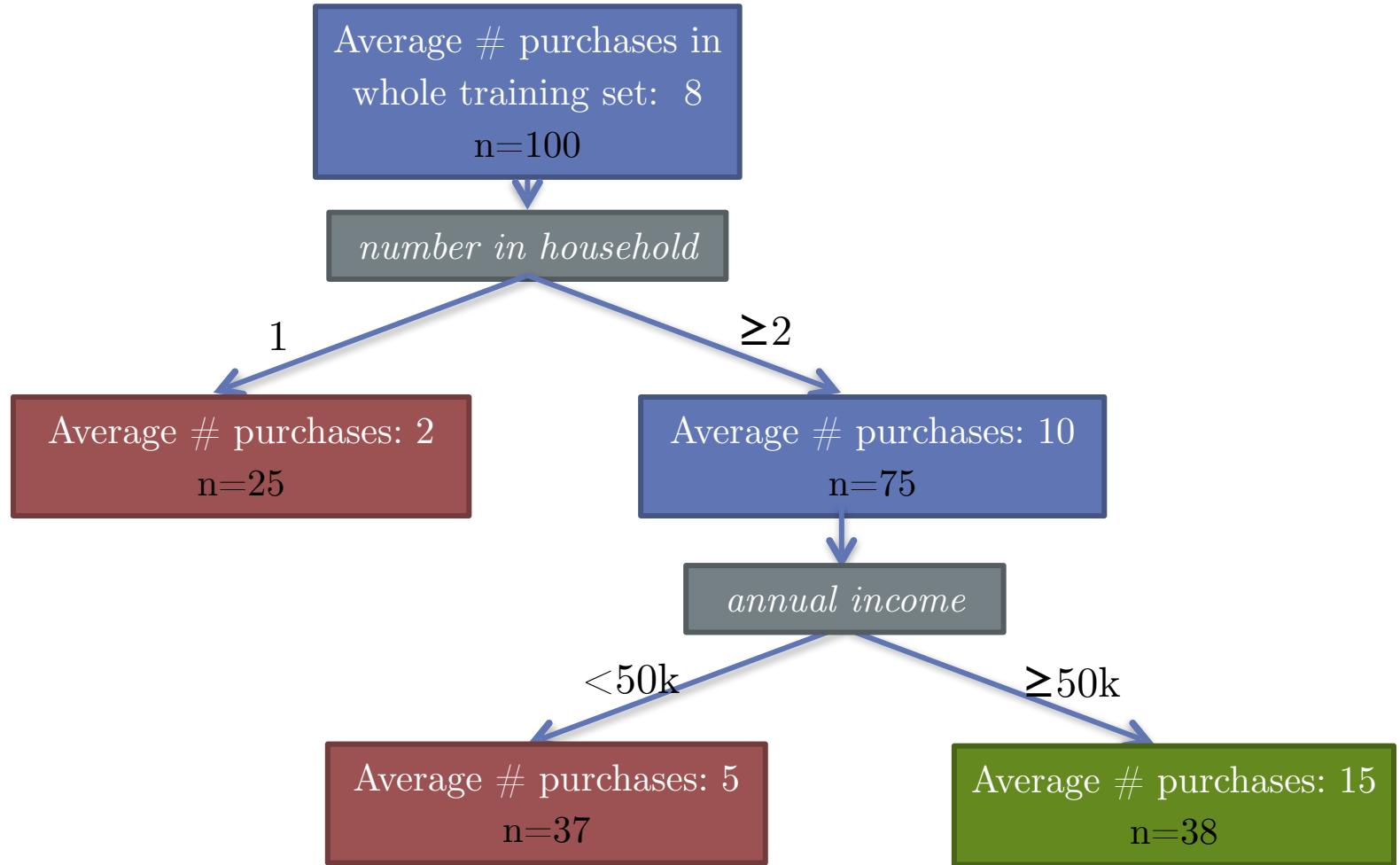
• • •

Same thing, but with **continuous target variables**

Regression Tree Model



Regression Tree Model Creation



Determining Splits

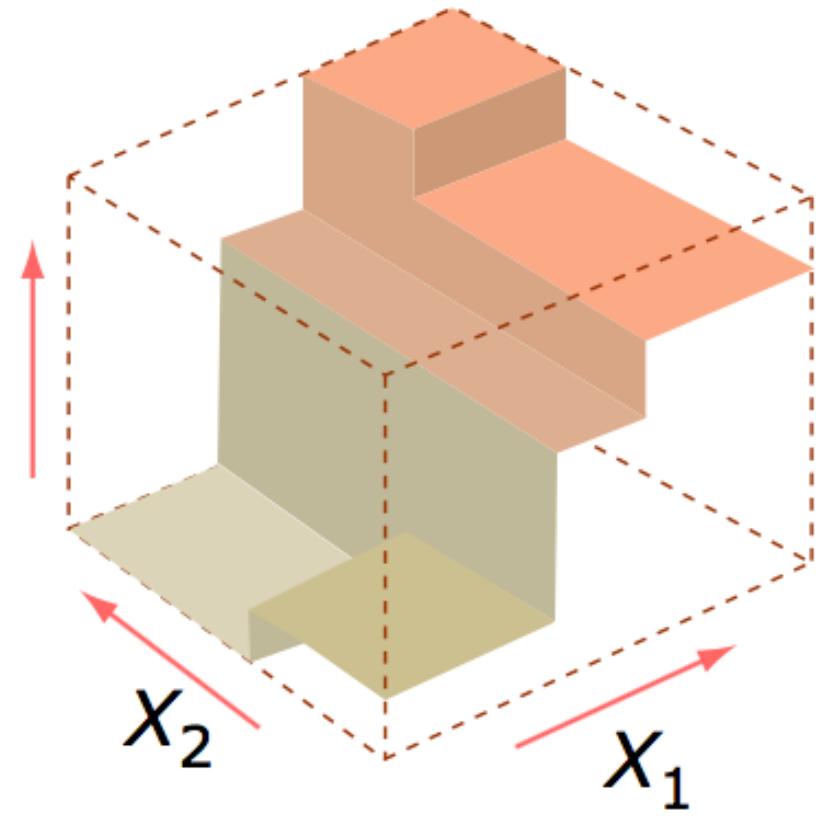
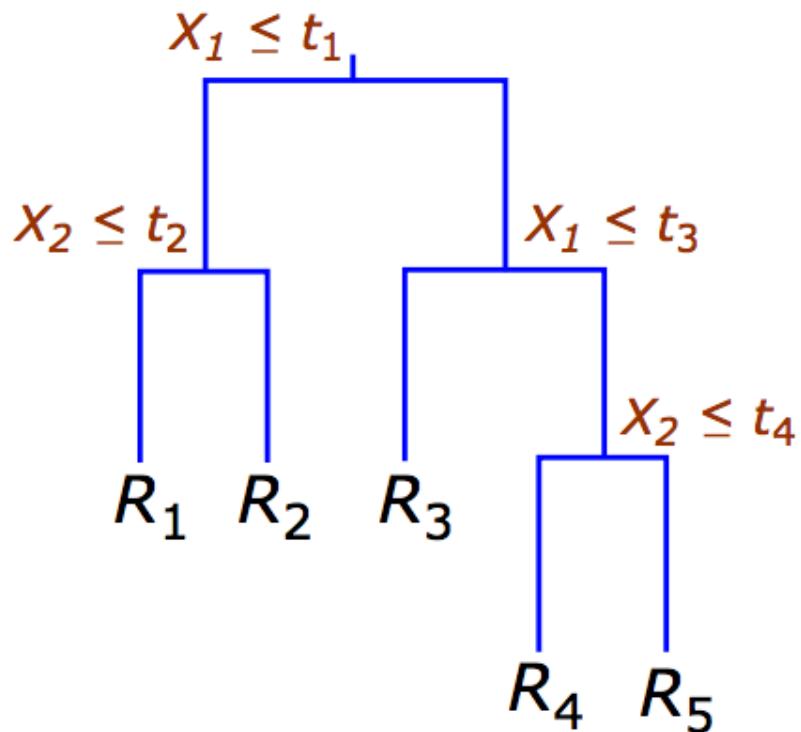
- Entropy/Gini no longer make sense for continuous target
- Instead:
 - Reduce *Average Squared Error* (i.e. variance since prediction is mean of observations in leaf)

$$\sum_{i=1}^{N_t} (y_i - \hat{y}_i)^2 = \sum_{i=1}^{N_t} (y_i - \bar{y}_i)^2 = \text{Var}(\mathbf{y}) \text{ within node}$$

- Or Maximize *logworth* using p-value from an F-test
 - Testing whether means (predicted value) of leaves is different
 - (Same as a t-test for difference of means in binary case)
 - Think ANOVA overall F-test: are any of these means different?

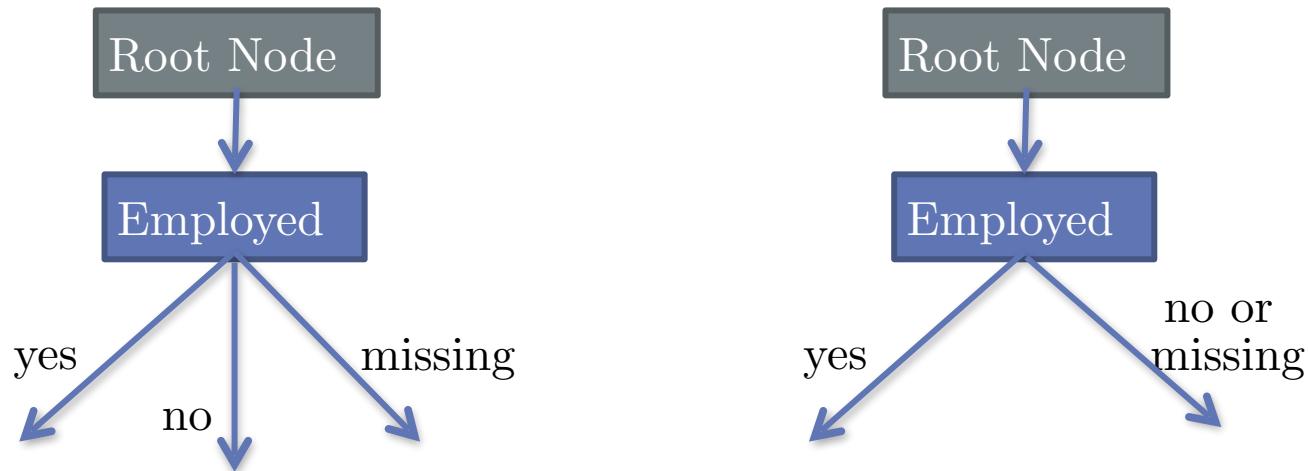
Regression Tree Response Surface

(Building with legos - no diagonals!)



Advantages of tree models

1. Explainability
2. Predicted probability/response has **meaning** in training set
3. Can handle missing values



Alternatively via **surrogate splits**: designate an alternative variable split if the given variable is missing. Surrogate splits are chosen in a way that they split the population in the most similar fashion to the current split (often use a highly correlated variable).

Advantages of tree models

1. Explainability
2. Predicted probability/response has **meaning** in training set
3. Can handle missing values
4. Can be used for **variable selection**
5. Great for **ensembles**
(basis for Random Forests and Gradient Boosting)
6. **No assumptions** to verify
7. Generally **immune to scale of input vars**/standardization
(less effort in data pre-processing)
8. Generally **immune to the effect of outliers** or high leverage observations

Disadvantages of tree models

1. **Simplistic** Regression/Decision Surface (non-smooth)
2. All variables forced to interact
 - a. Only the top split acts independently
 - b. Inefficient
3. **Greedy** Algorithms
 - a. Struggle in the presence of many variables
 - b. Cannot return the globally optimal tree
4. Can be **unstable** (sensitive to small changes in input) - both when training the model *and* when making predictions.
(think: sides of ‘lego buildings’ on the response surface)

Viya Demo 2

• • •

TelcoChurn using Tasks in SAS Studio

ANALYTICS LIFE CYCLE

Manage Data

Prepare Data

Explore and Visualize

Build Models

Manage Models

Share and Collaborate

Develop SAS Code

Tasks

Filter

My Tasks

SAS Tasks

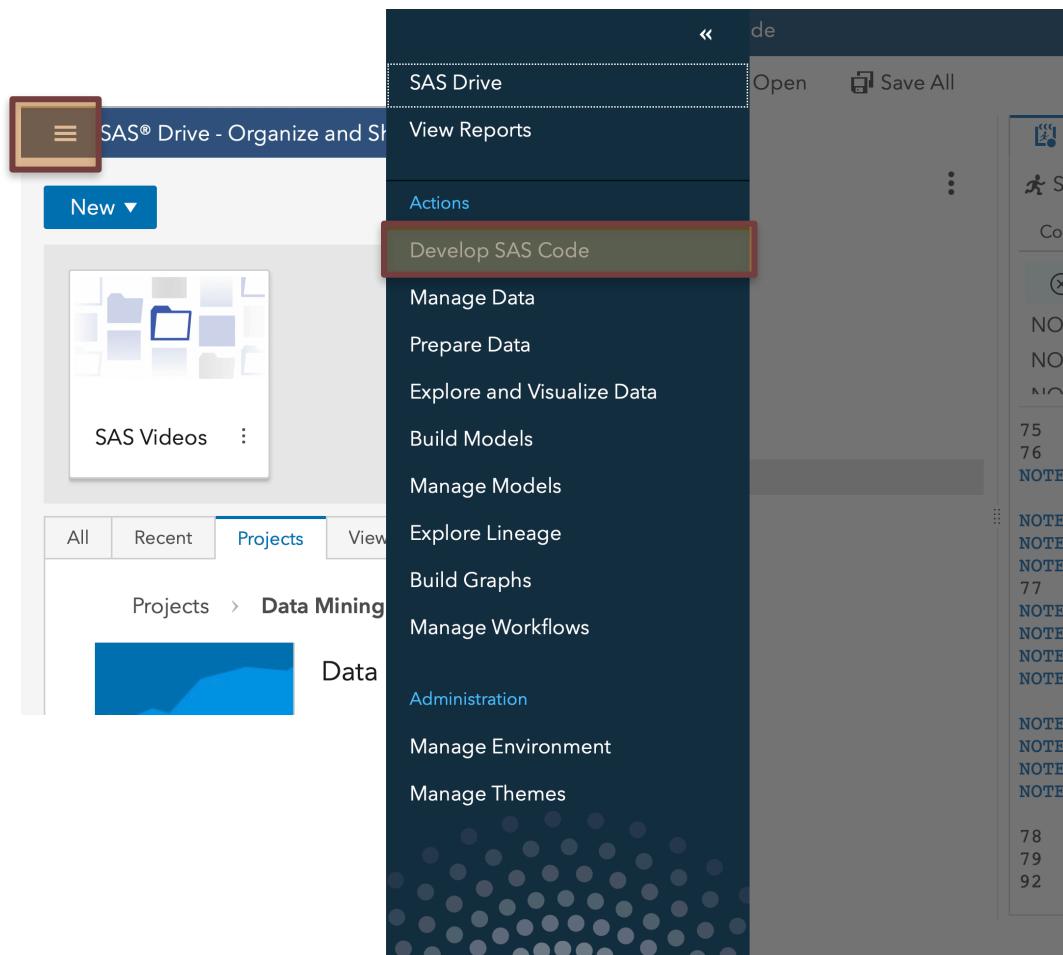
- ▶ Prepare Data
- ▶ Visualize Data
- ▶ Statistics
- ▶ Econometrics
- ▶ Forecasting
- ▶ Optimization and Network Analysis
- ▶ Statistical Process Control
- ▶ SAS Viya Cloud Analytic Services
- ▶ SAS Viya Prepare and Explore Data
- ▶ SAS Viya Evaluate and Implement Models
- ▶ SAS Viya Statistics
 - Clustering
 - Principal Component Analysis
 - Linear Regression
 - Logistic Regression
 - Generalized Linear Models
 - Partial Least Squares Regression
 - Quantile Regression
 - Decision Tree
- ▶ SAS Viya Machine Learning
- ▶ SAS Viya Econometrics
- ▶ SAS Viya Forecasting
- ▶ SAS Viya Text Analytics
- ▶ SAS Viya Optimization and Network Analysis

Viya Demo 3

• • •

Breast Cancer Malignancy

Viya Demo



Submit Code:

```
cas;  
caslib _all_ assign;
```

You will repeat this step
EVERY time you use
Viya to load the Public
library!

Identifying Malignant Tumors

The screenshot shows the SAS Visual Analytics interface. At the top left is the 'SAS® Drive - Organize and Share Content' bar with a 'New ▾' button. Below it is a sidebar with 'SAS Videos' and a 'Projects' tab selected. A dropdown menu is open over the 'Projects' tab, listing options: 'SAS Drive', 'View Reports', 'Actions', 'Develop SAS Code' (which is highlighted in blue), 'Manage Data', 'Prepare Data', and 'Explore and Visualize Data' (which is highlighted in gold). To the right of the sidebar is a main area titled 'Welcome to SAS Visual Analytics' with a sub-instruction 'Select an option to get started:' followed by three icons: a grid labeled 'Data', a chart labeled 'New', and a folder labeled 'Open'. At the bottom of the main area is a checkbox with the text 'Make this selection the default'. Below this is a section titled 'Open Data Source' with tabs for 'Available', 'Data Sources', and 'Import'. The 'Available' tab is selected, showing a list of data sources. The first item in the list is 'BANK' (09/20/19 11:30 AM • slrace), which is highlighted in orange. The second item is 'BREAST_CANCER' (09/20/19 11:30 AM • slrace), also highlighted in orange. The third item is 'CARS_TRAIN' (08/29/19 09:22 AM • slrace).

Welcome to SAS Visual Analytics

Select an option to get started:



Make this selection the default

Open Data Source

Available Data Sources Import

Filter



BANK
09/20/19 11:30 AM • slrace

BREAST_CANCER
09/20/19 11:30 AM • slrace

CARS_TRAIN
08/29/19 09:22 AM • slrace

The screenshot shows a user interface for a data visualization or machine learning application. The left sidebar features three main sections: 'Data' (selected), 'Objects' (unselected), and 'Outline' (unselected). The main area is titled 'Data' and contains a search bar with the text 'BREAST_CANCER'. Below the search bar is a 'Filter' button with a magnifying glass icon. A large orange callout box highlights a dropdown menu item labeled '0(benign) 1(malignant)'. This menu is part of a larger panel with tabs for 'Category' (selected), 'Measure' (highlighted in blue), and 'Format'. The 'Format' tab has a dropdown arrow and a small edit icon. At the bottom, the text 'Format:' is displayed.

Data

BREAST_CANCER

-
-
-
-
-
-
-
-
-

Name:

Based on:
 Data item Sampling

Sampling method:

Number of partitions:

Training partition sampling percentage: *

Random number seed

Random seed: *

Change target attribute to categorical variable (split into training/validation)

Objects

Filter

▼ SAS Visual Statistics

Cluster

Decision Tree

Generalized Additive Model

Data Roles

Decision Tree - 0(benign) 1(malignant) 1 ▾

▼ Response

0(benign) 1(malignant)

▼ Predictors

- ❖ Marginal Adhesion
- ❖ Mitoses
- ❖ Normal Nucleoli
- ❖ Single Epithelial Cell Size
- ❖ Uniformity of Cell Shape
- ❖ Uniformity of Cell Size
- ❖ Bare Nuclei
- ❖ Bland Chromatin
- ❖ Clump Thickness
- ✚ Add

▼ Partition ID

Validation

Options
Roles

Actions

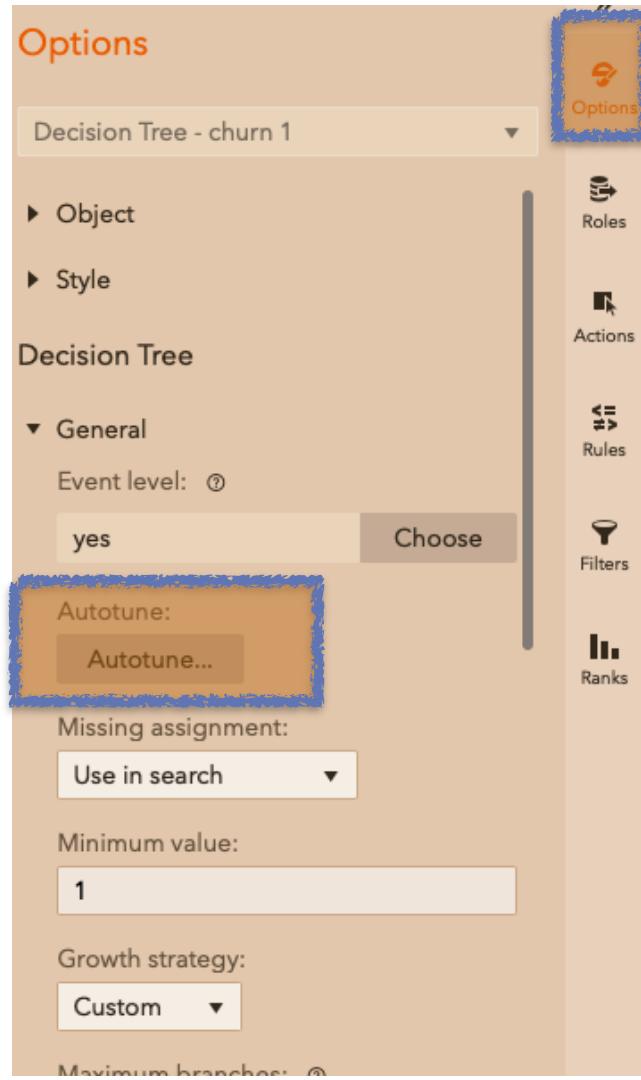
Rules

Filters

Ranks

Create a decision tree and set the

Autotune Function



Stack Display

Options

Decision Tree - churn 1

- Rapid growth
- Prune with validation data
- Pruning: 75%
- Reuse predictors

Assessment

Model Display

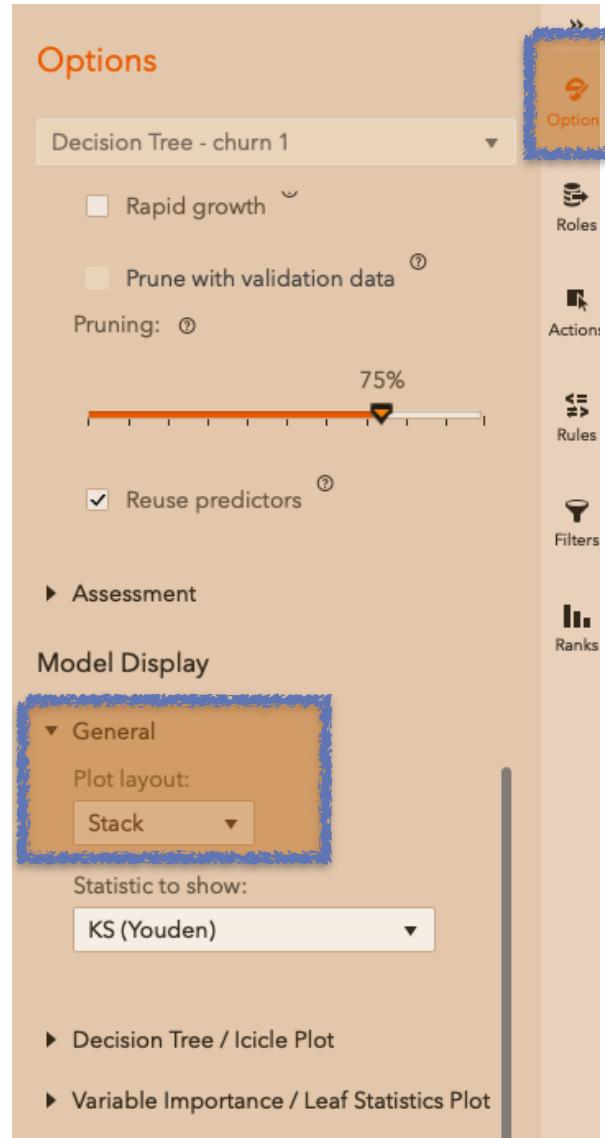
General

Plot layout: Stack

Statistic to show: KS (Youden)

Decision Tree / Icicle Plot

Variable Importance / Leaf Statistics Plot



[«](#)

Decision Tree

Icicle

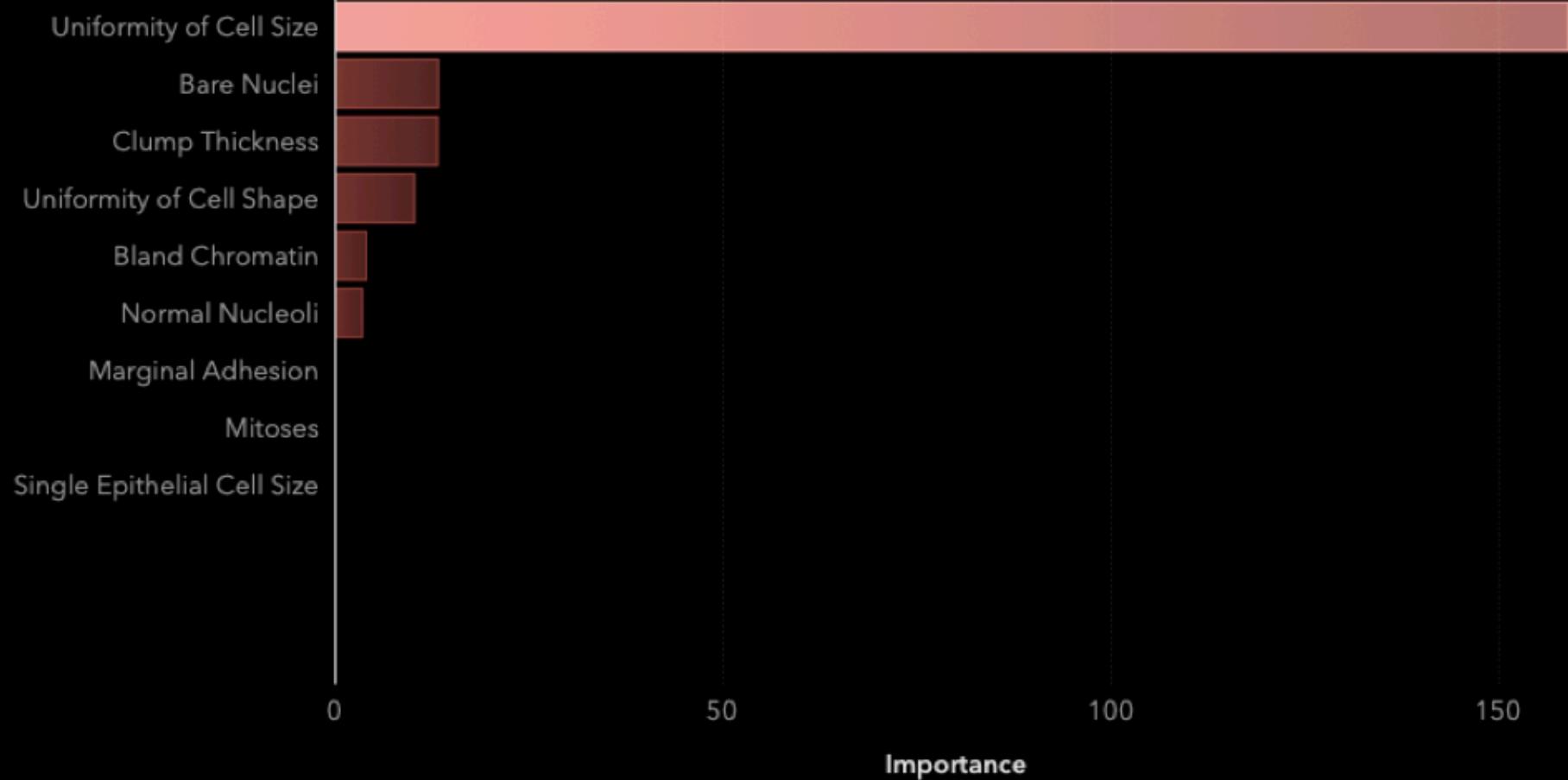
Variable Importance

Assessment

[»](#)

⋮

Variable Importance

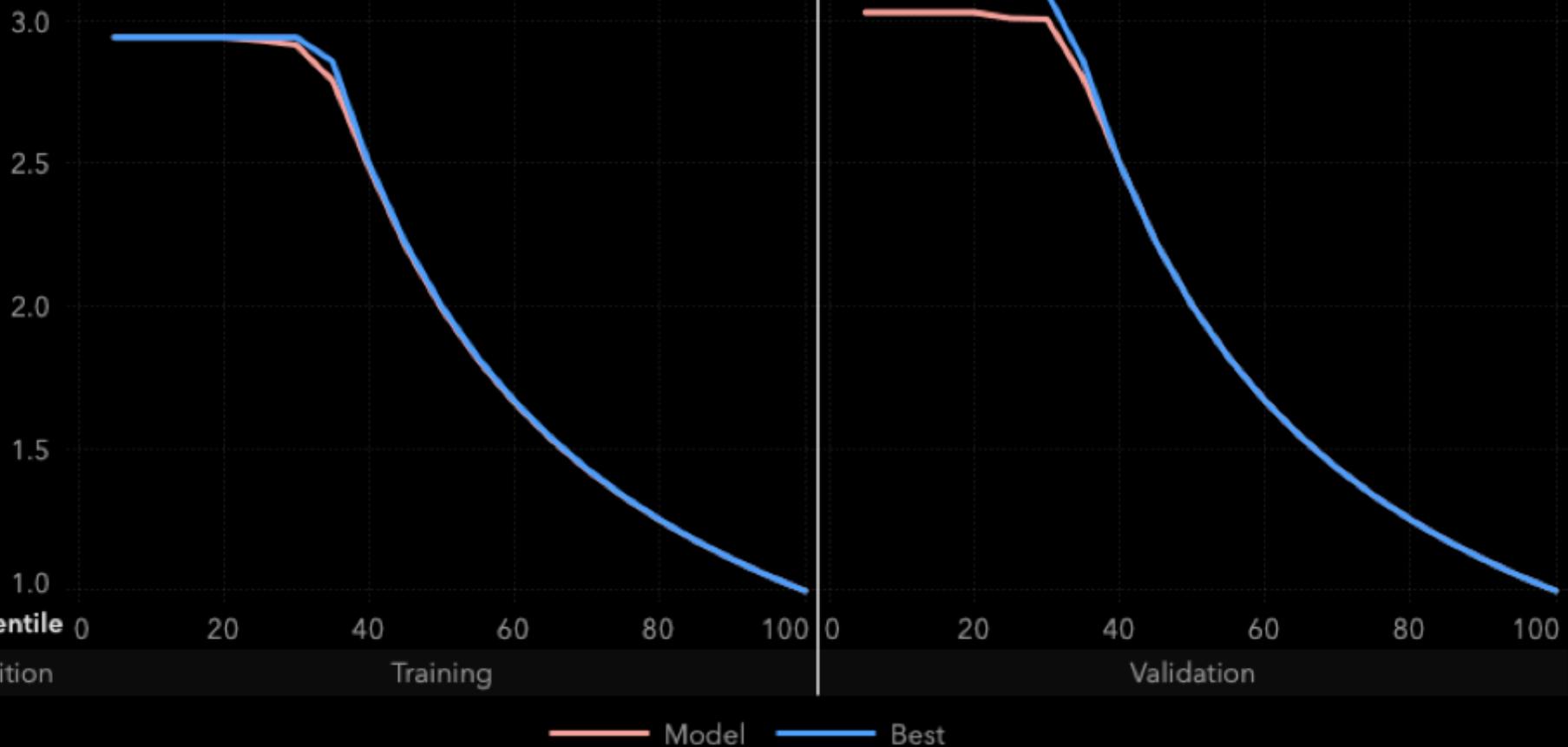


Decision Tree Icicle Variable Importance

Assessment

Lift

Cumulative Lift

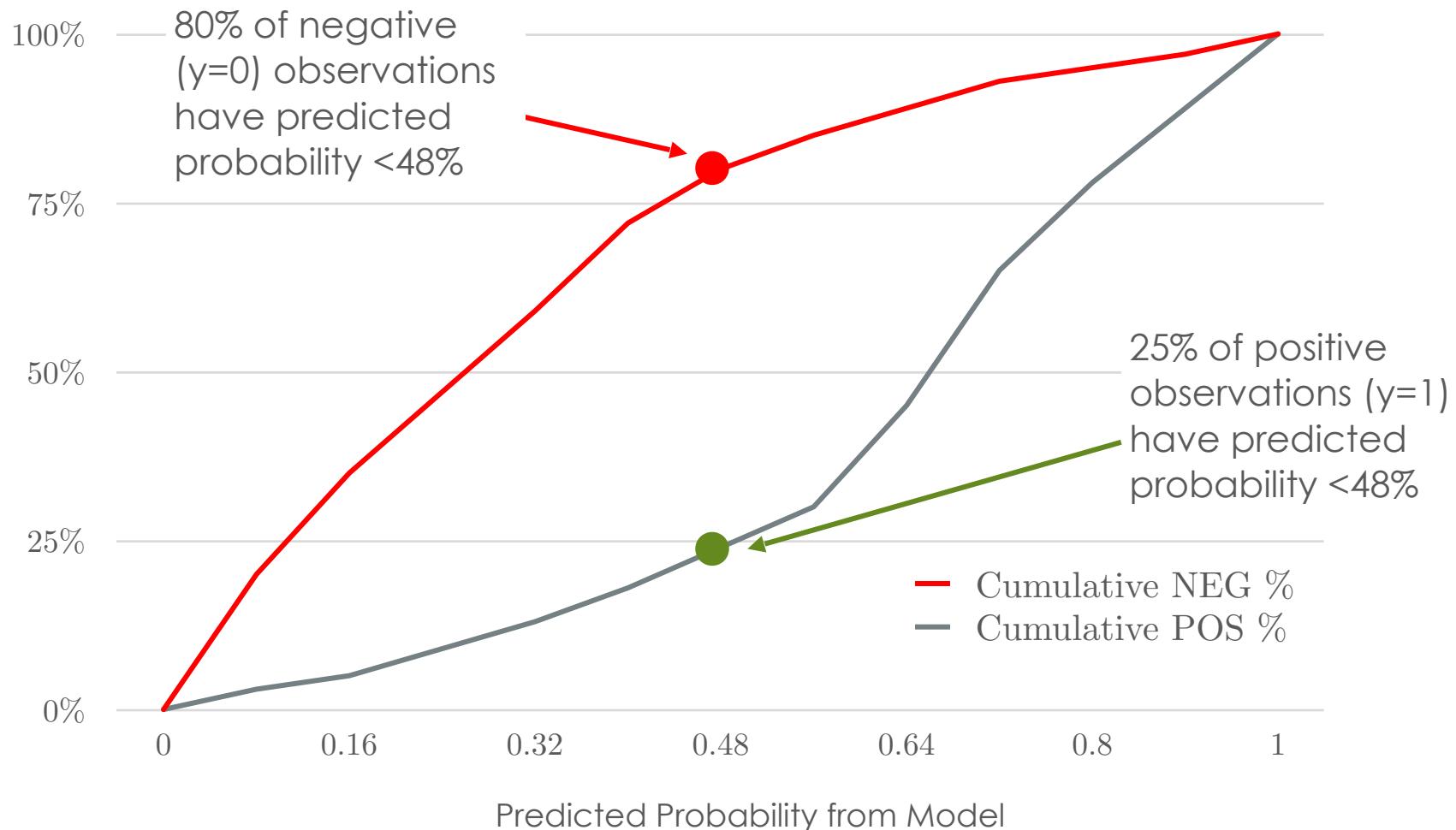


Additional Reference Slides

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The K-S Statistic

Kolmogorov-Smirnov (KS) Statistic



Kolmogorov-Smirnov (KS) Statistic

