# Tree pruning and bagging **STAT 4710**

October 31, 2023

### Where we are

Unit 1: R for data mining
Unit 2: Prediction fundamentals
Unit 3: Regression-based methods
Unit 4: Tree-based methods
Unit 5: Deep learning

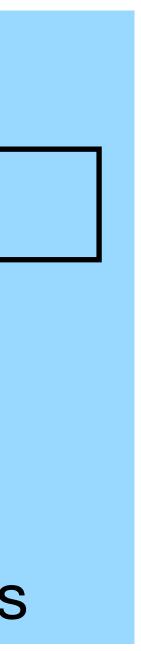
Lecture 1: Growing decision trees

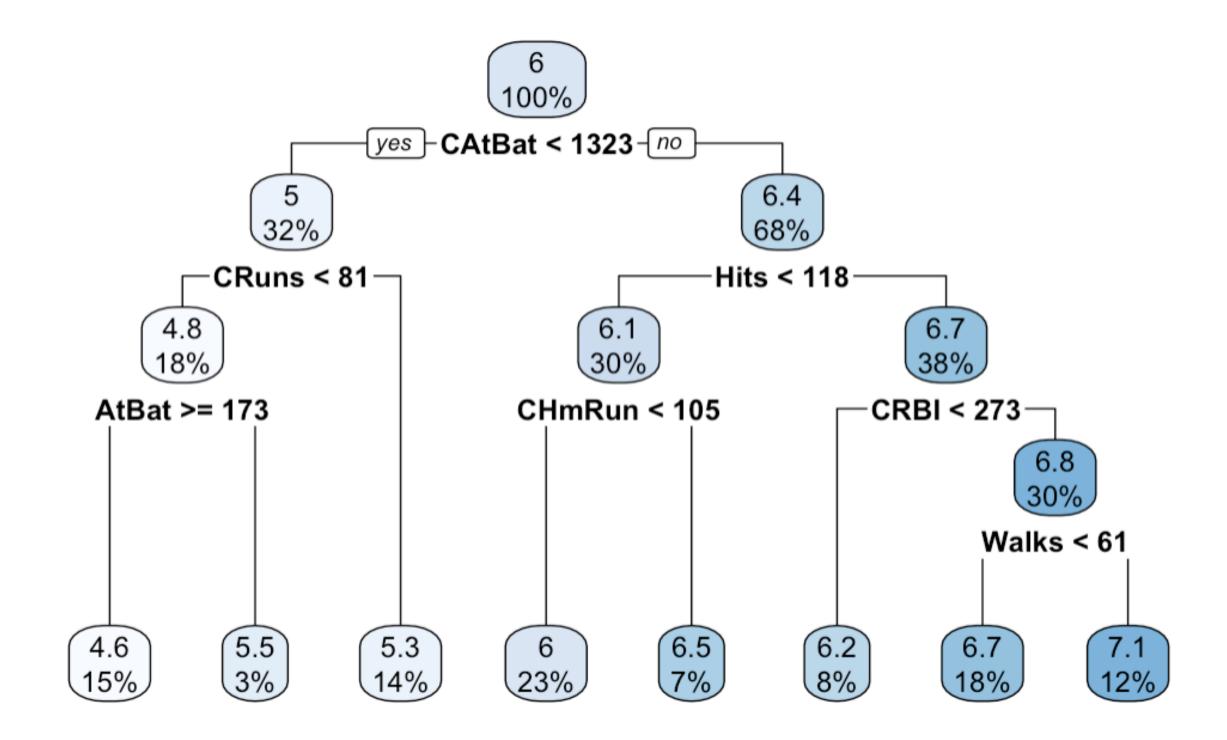
**Lecture 2:** Tree pruning and bagging

Lecture 3: Random forests

Lecture 4: Boosting

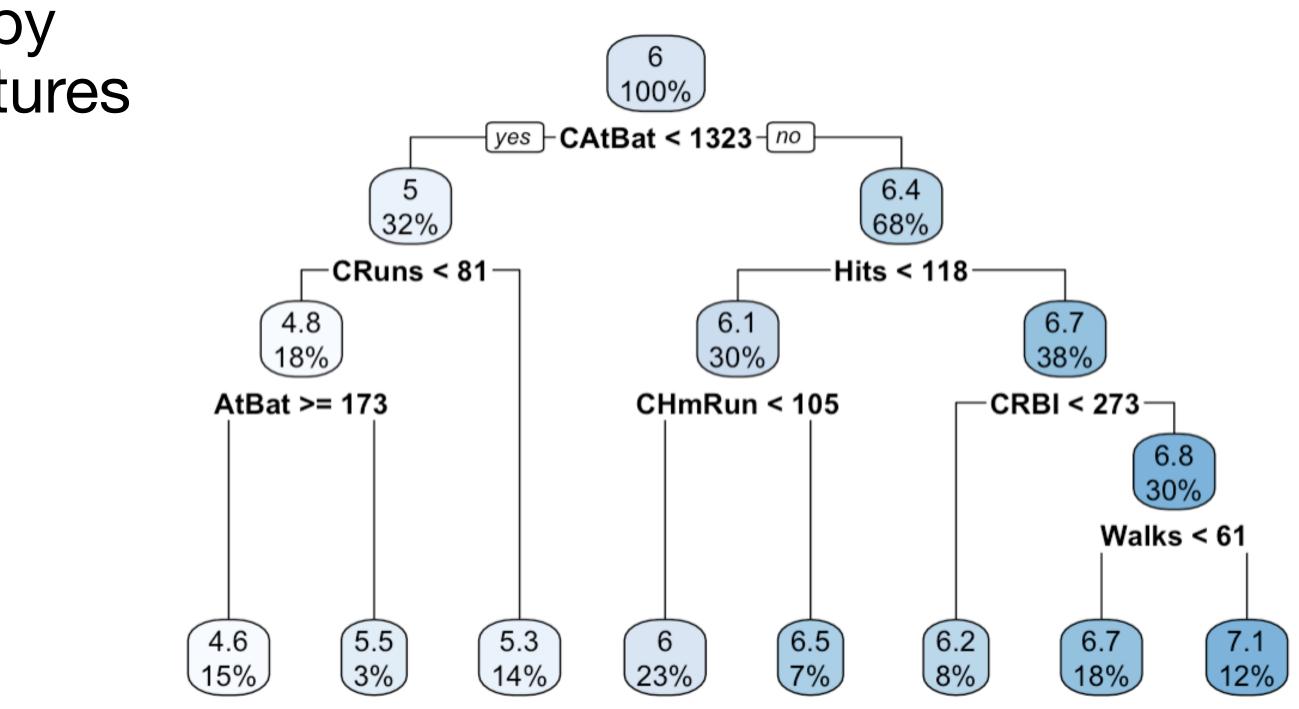
Lecture 5: Unit review and quiz in class





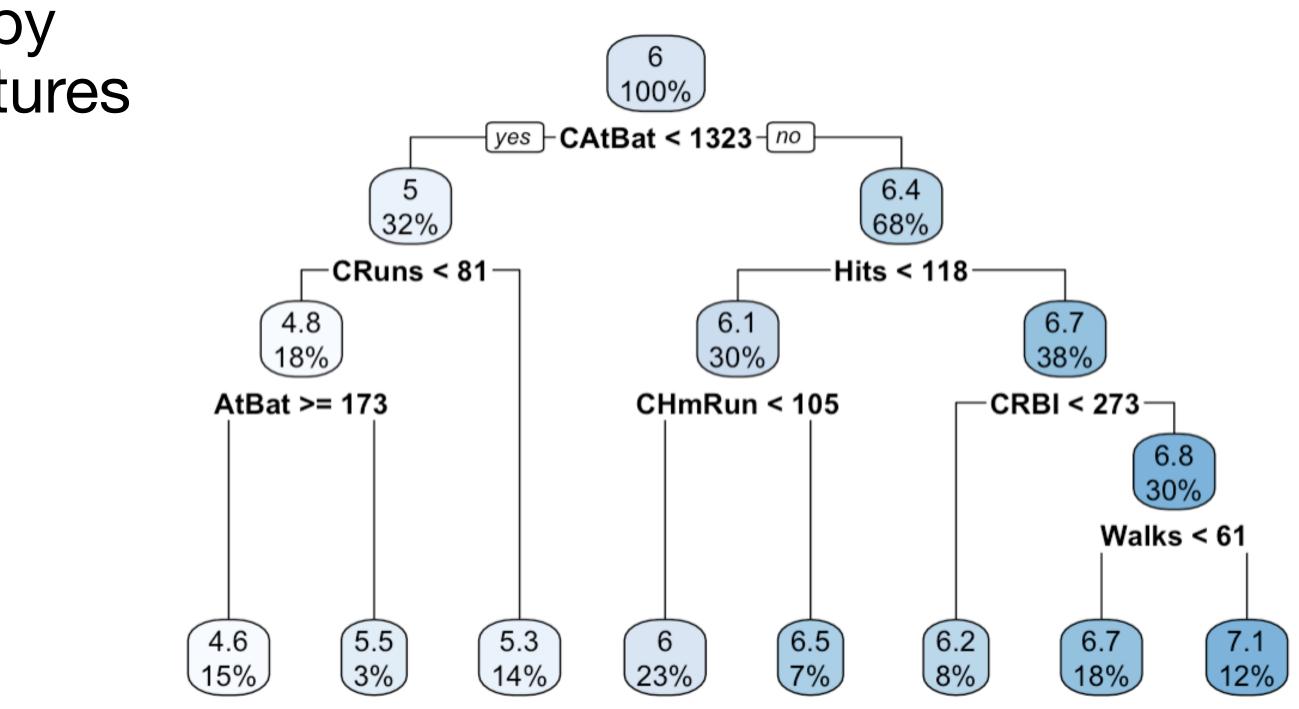
 Create a partition of feature space by recursively splitting on different features





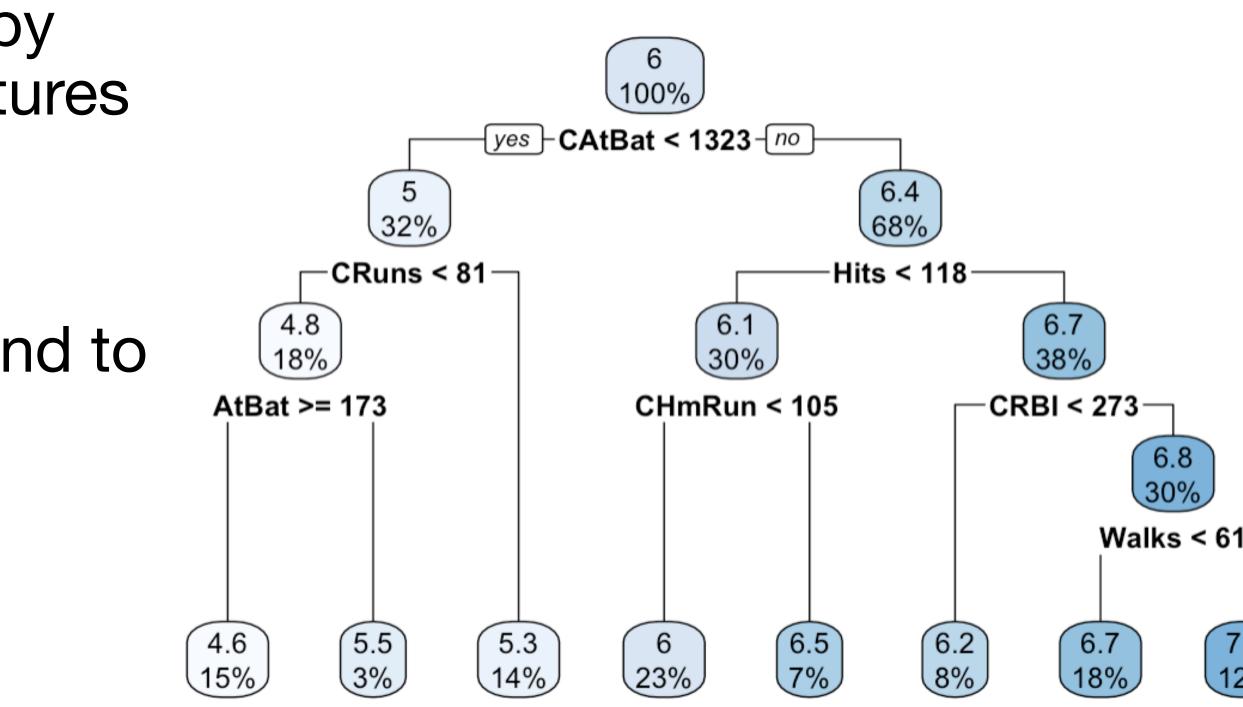
- Create a partition of feature space by recursively splitting on different features
- Regression and classification trees

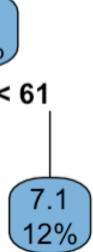




- Create a partition of feature space by recursively splitting on different features
- **Regression and classification trees**
- Terminal nodes in the tree correspond to the rectangles in the partition

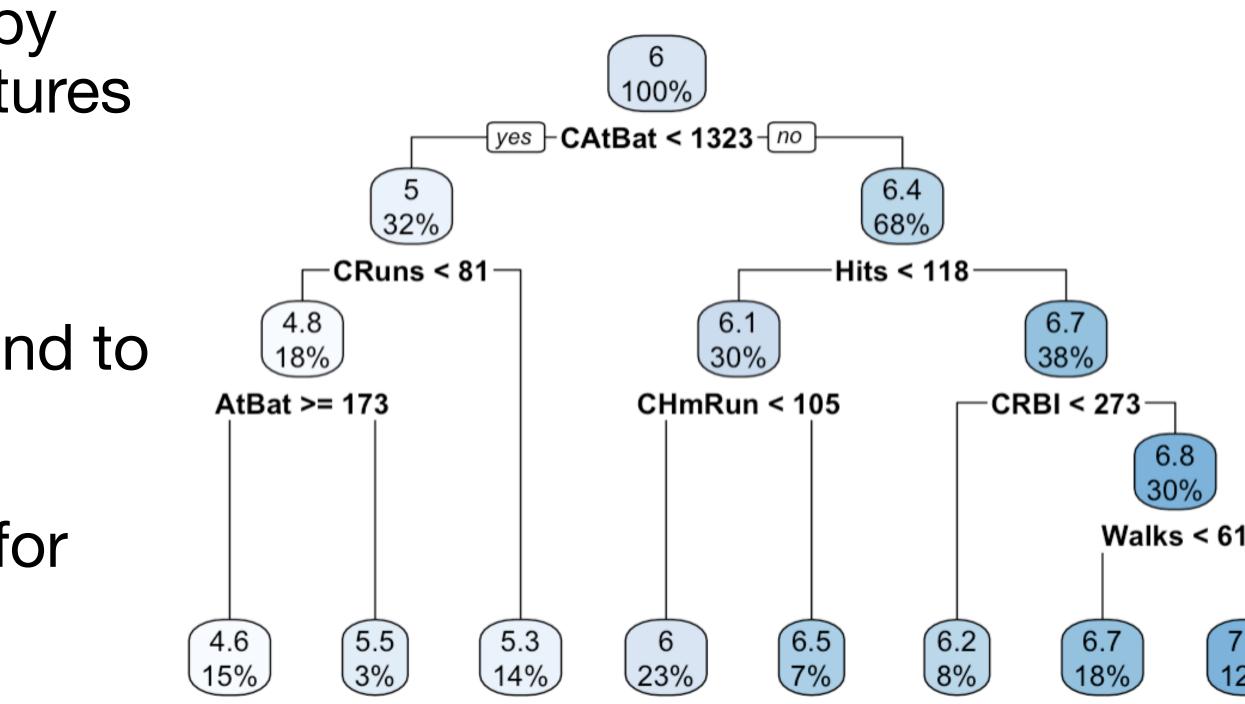






- Create a partition of feature space by recursively splitting on different features
- Regression and classification trees
- Terminal nodes in the tree correspond to the rectangles in the partition
- Predict a single number (category) for each terminal node in a regression (classification) tree







- if there are as many terminal nodes as training points, training error = 0
- If there is just one terminal node, we are fitting a constant model

The more terminal nodes (regions), the more flexibly the tree fits training data:

- if there are as many terminal nodes as training points, training error = 0
- If there is just one terminal node, we are fitting a constant model

complexity.

The more terminal nodes (regions), the more flexibly the tree fits training data:

As with any prediction method, there is a bias-variance tradeoff based on model

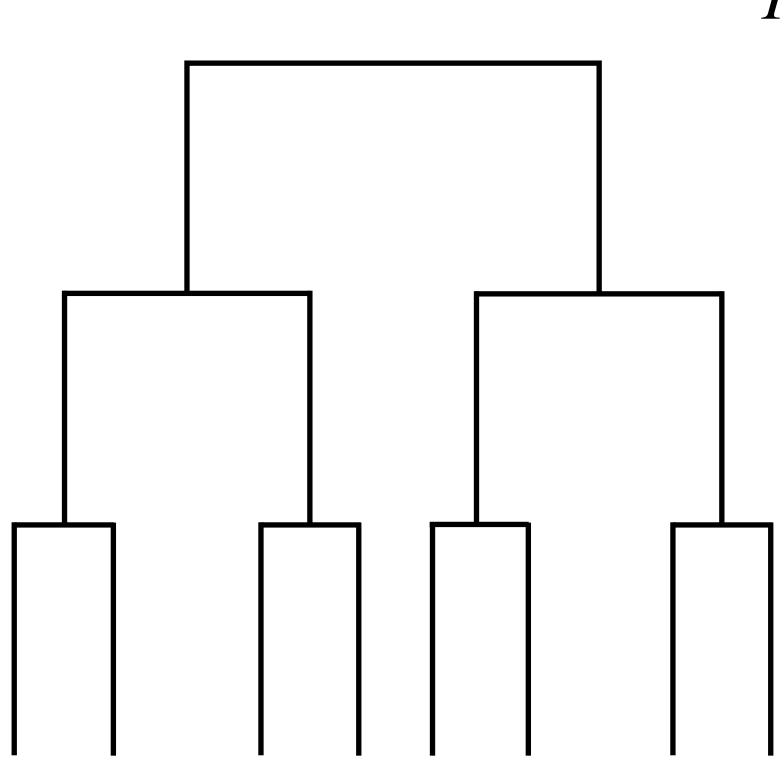
- if there are as many terminal nodes as training points, training error = 0
- If there is just one terminal node, we are fitting a constant model

complexity.

How to choose the best model complexity? Cross-validation.

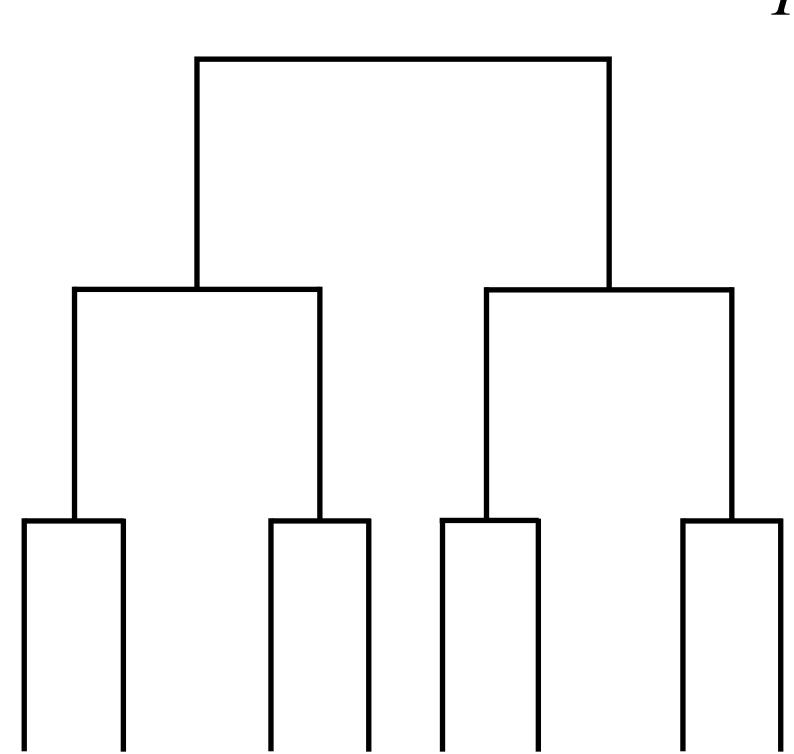
The more terminal nodes (regions), the more flexibly the tree fits training data:

- As with any prediction method, there is a bias-variance tradeoff based on model



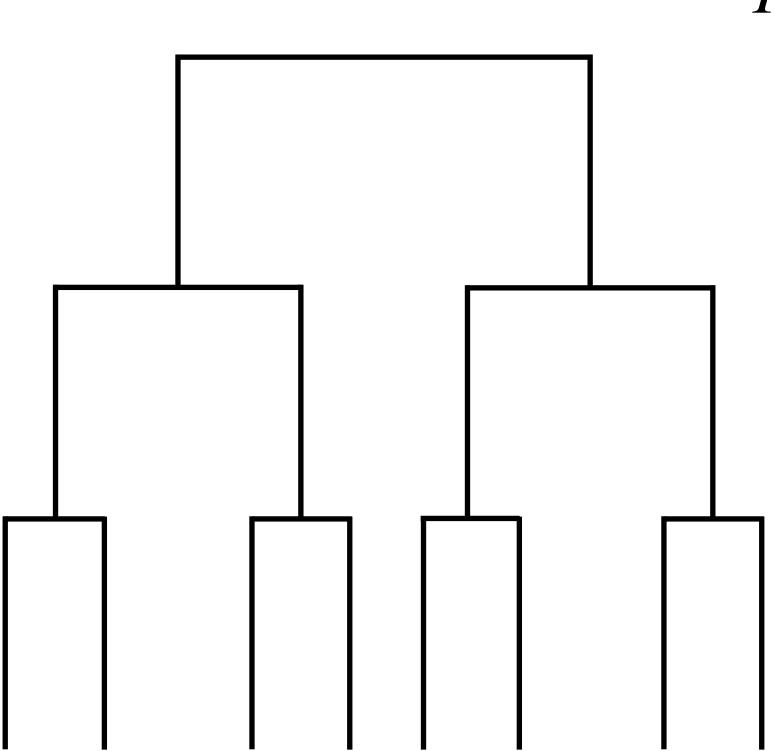


 First grow out our tree about as far as we can to obtain a big tree  $T_0$ .



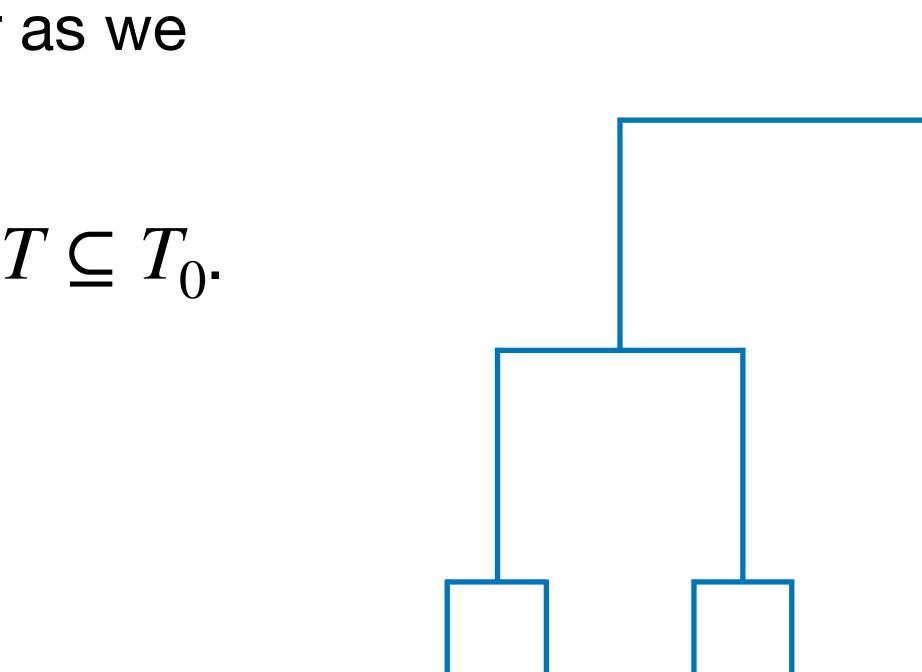


- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .



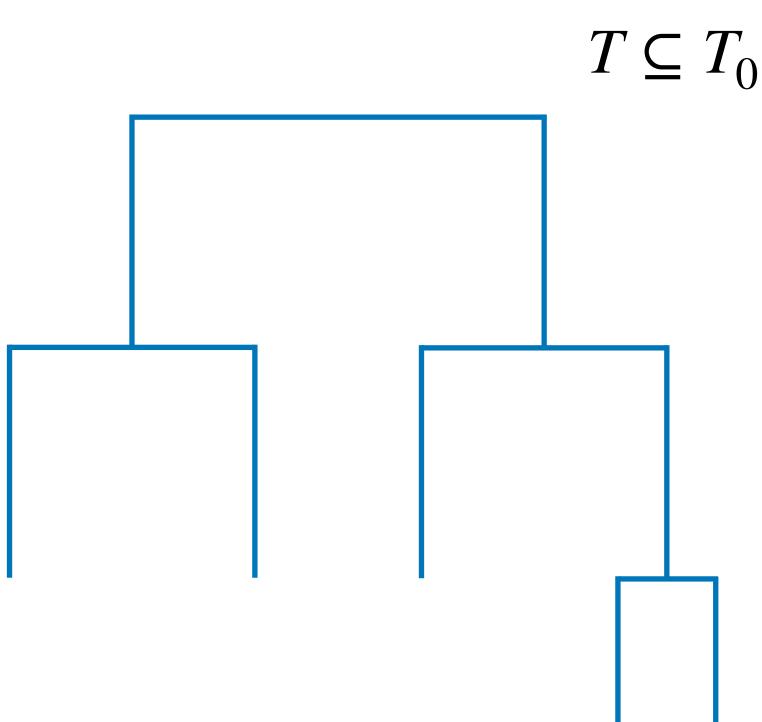


- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .





- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .





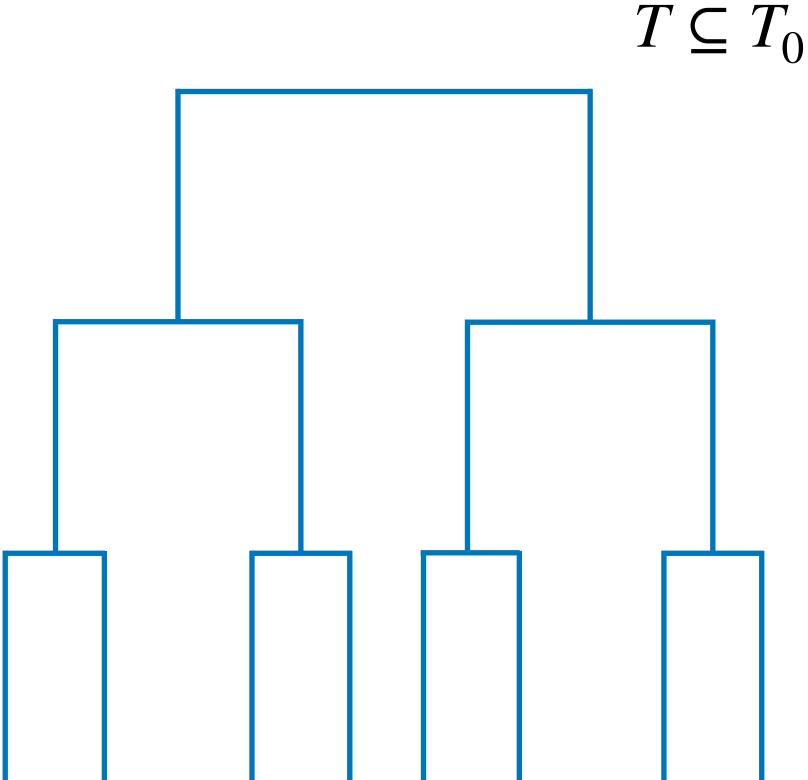
- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .

 $T \subseteq T_0$ 



- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .

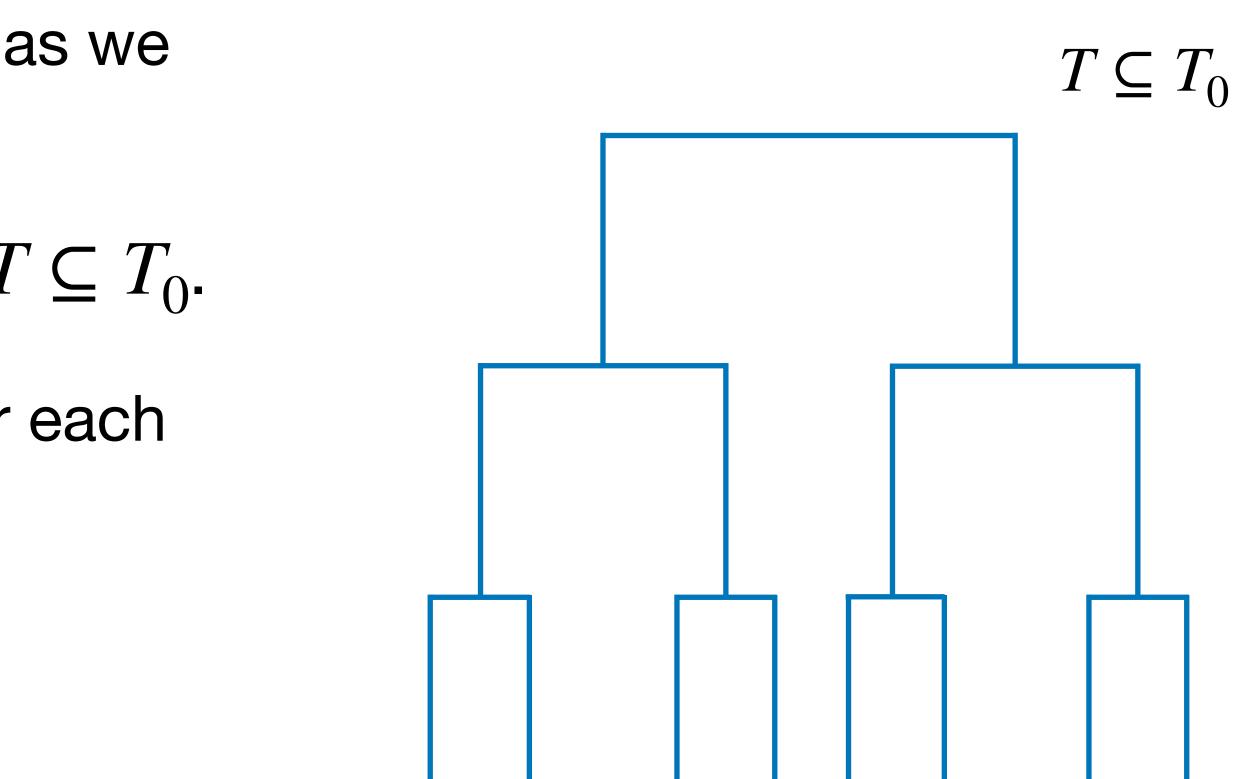






- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .

Note: There are several subtrees T for each complexity value.

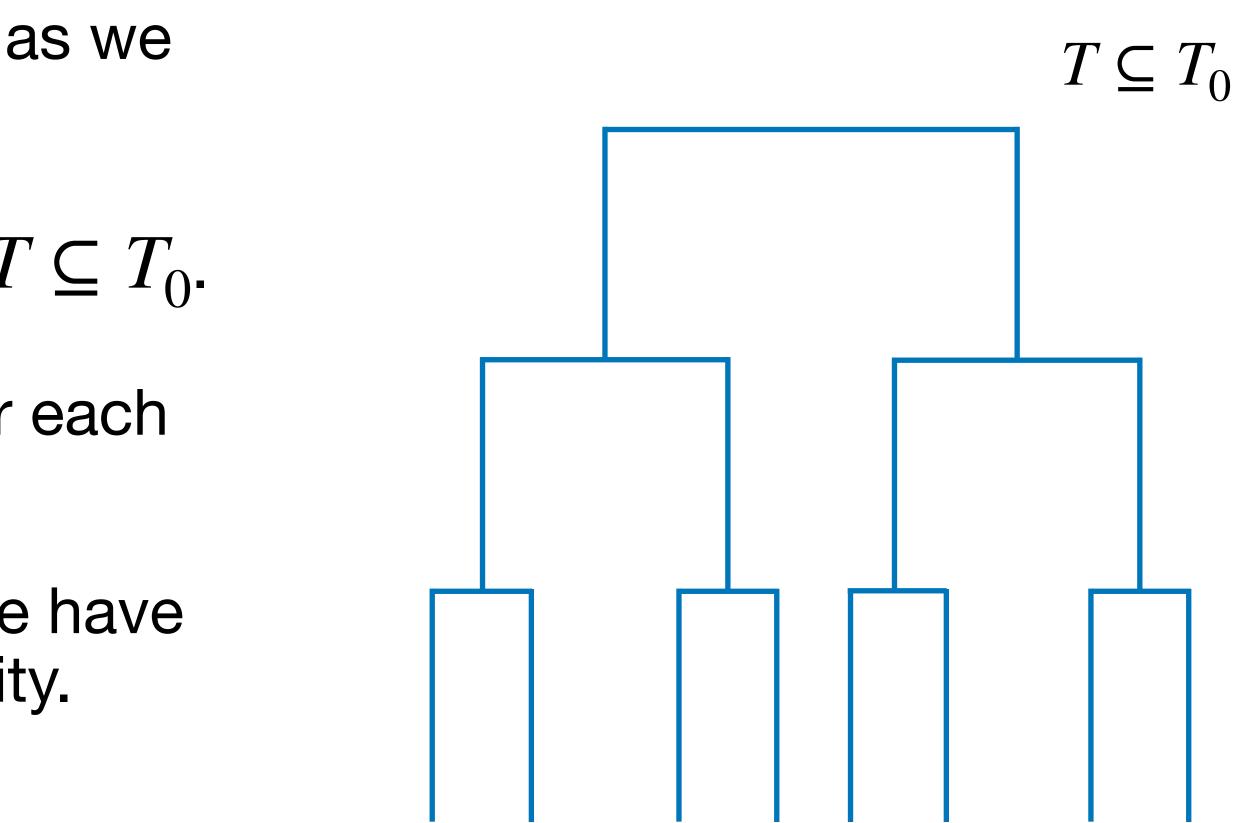




- First grow out our tree about as far as we can to obtain a big tree  $T_0$ .
- We can then consider any subtree  $T \subseteq T_0$ .

Note: There are several subtrees T for each complexity value.

In other model selection scenarios, we have just had one model for each complexity.







# $T \subseteq T_0$

Let |T| be number of terminal nodes in tree T. Fixing some  $\alpha \ge 0$ , consider  $T_{\alpha} = \arg\min \{ \operatorname{RSS}(T) + \alpha | T | \}.$ 



 $T \subseteq T_0$ 

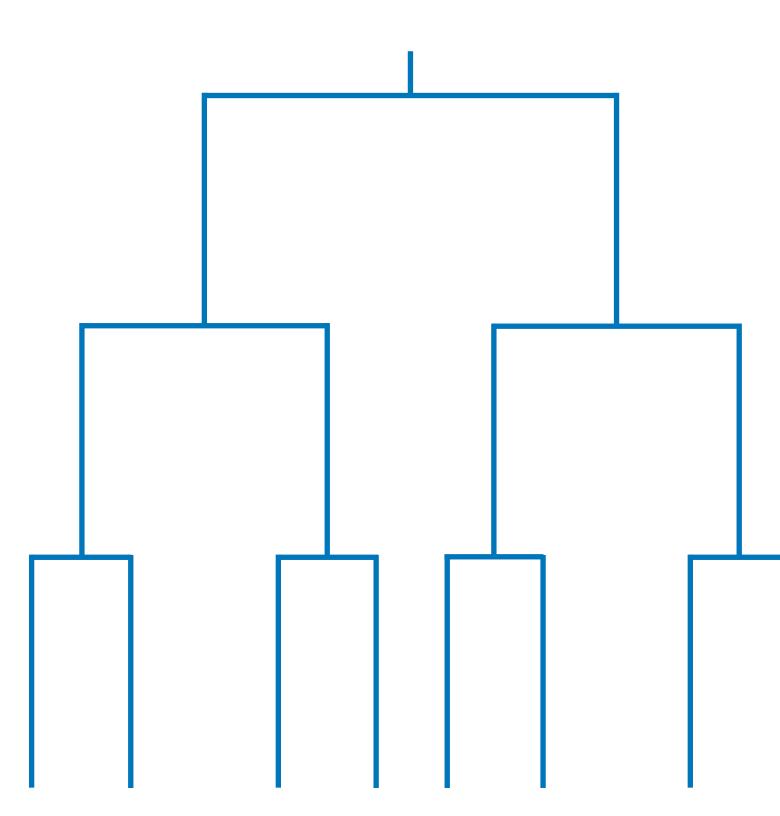
- Let |T| be number of terminal nodes in tree T. Fixing some  $\alpha \ge 0$ , consider  $T_{\alpha} = \arg\min \{ \operatorname{RSS}(T) + \alpha |T| \}.$
- Like lasso, varying  $\alpha$  leads to sequence of trees; higher  $\alpha$  leads to smaller trees.



 $T \subseteq T_0$ 

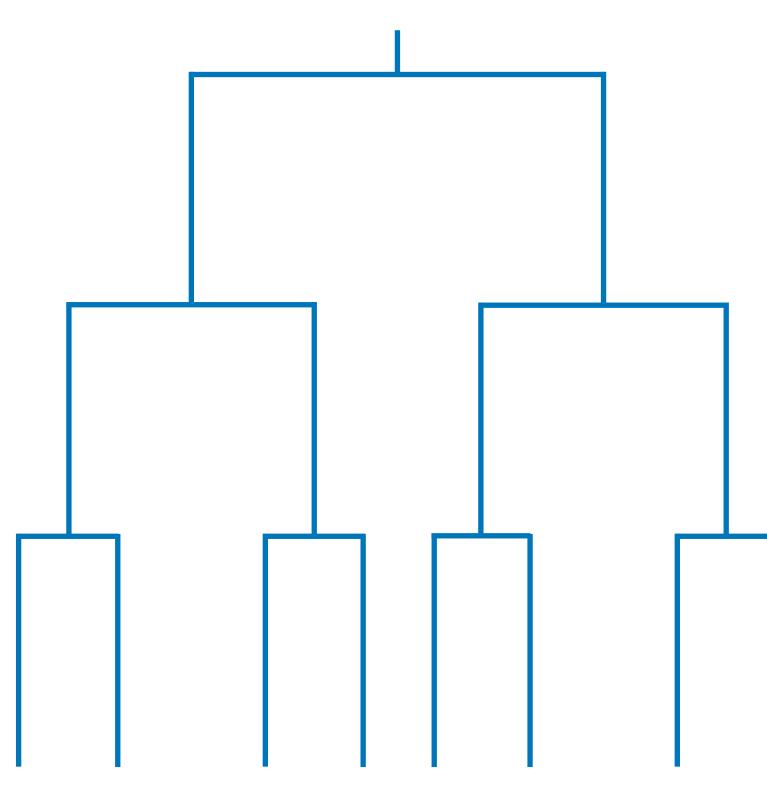
- Let |T| be number of terminal nodes in tree T. Fixing some  $\alpha \ge 0$ , consider  $T_{\alpha} = \arg\min \{ \operatorname{RSS}(T) + \alpha |T| \}.$
- Like lasso, varying  $\alpha$  leads to sequence of trees; higher  $\alpha$  leads to smaller trees.
- Unlike lasso, discrete set of  $\alpha$  values gives all possible solutions as  $\alpha$  varies.







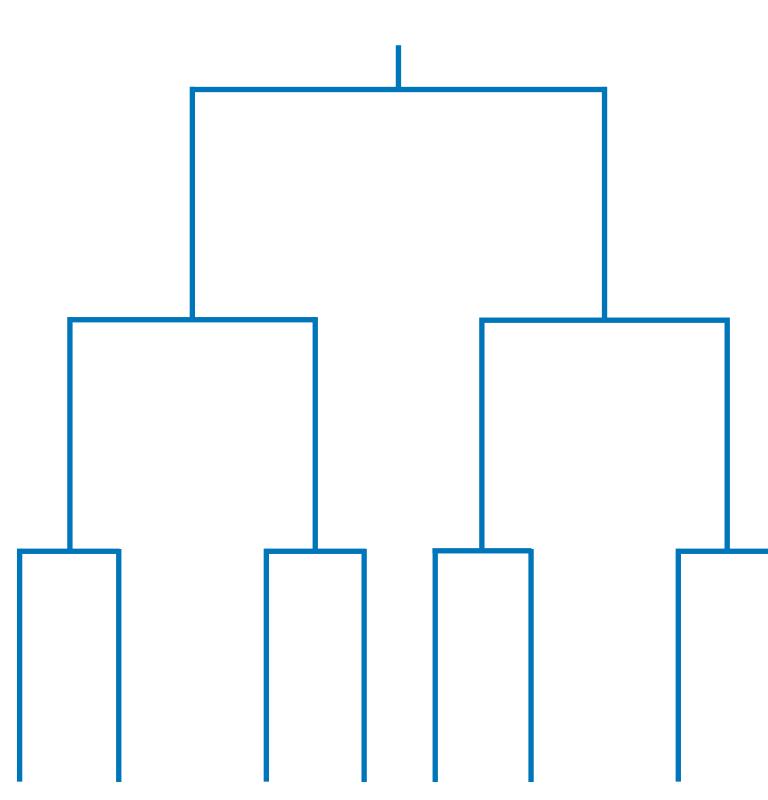
Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.





Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

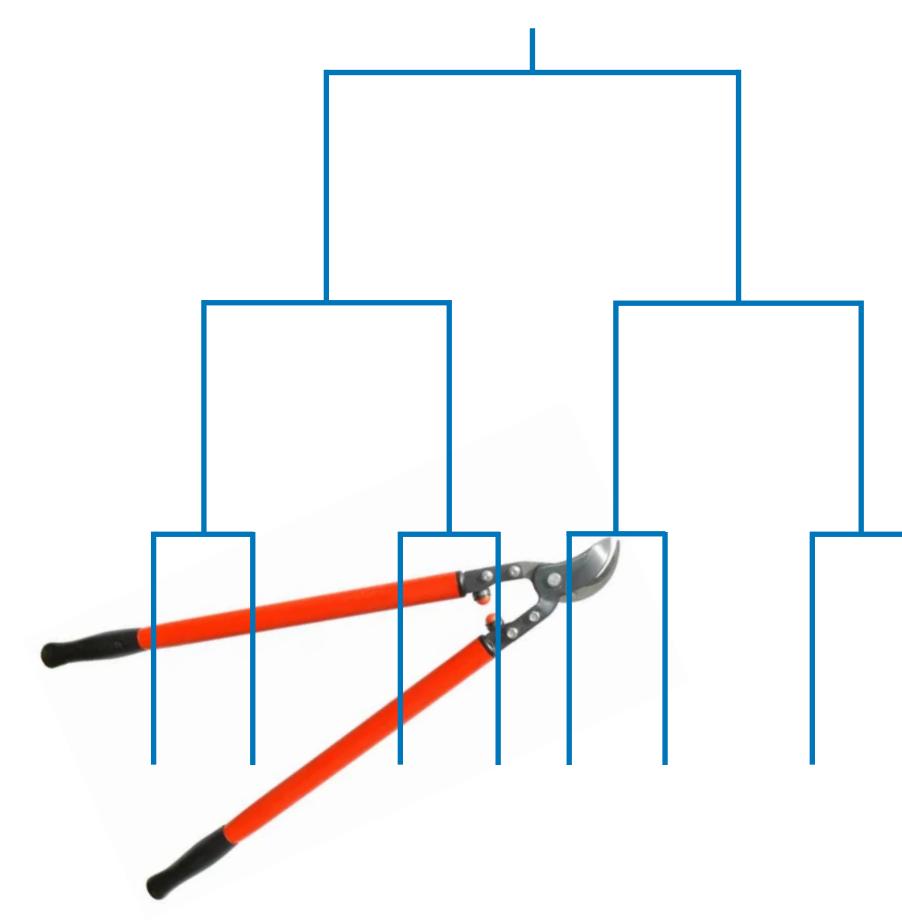
Idea: Recursively prune "weakest link" splits.





Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

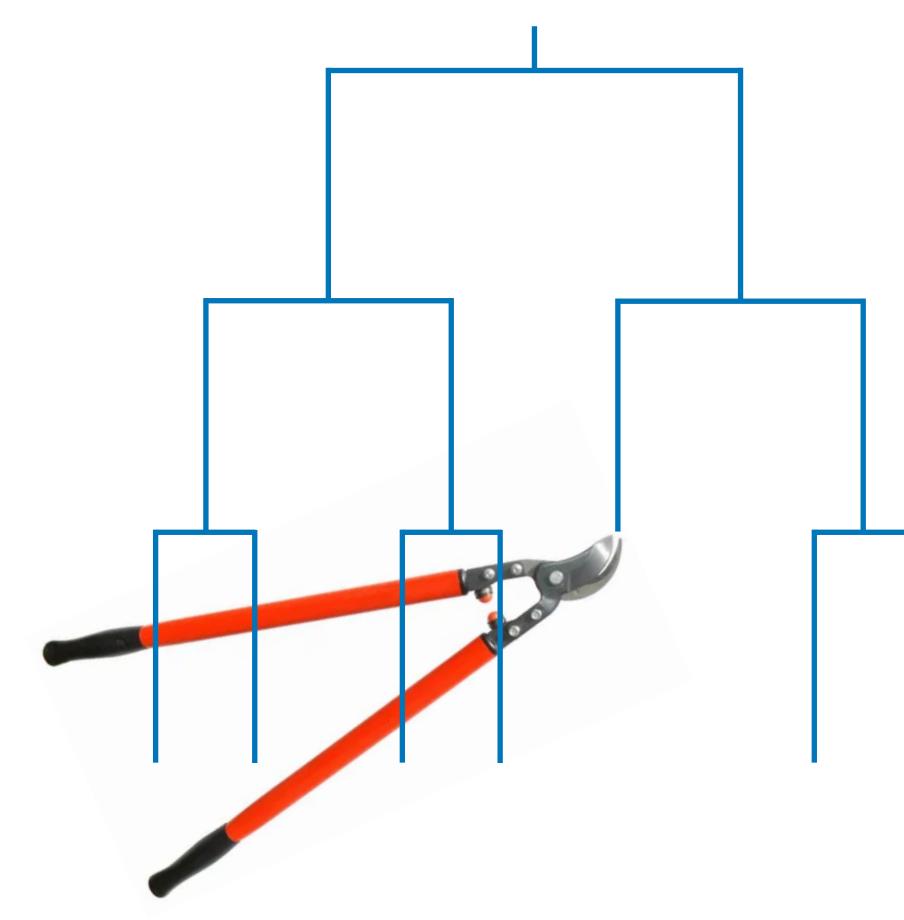
Idea: Recursively prune "weakest link" splits.





Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

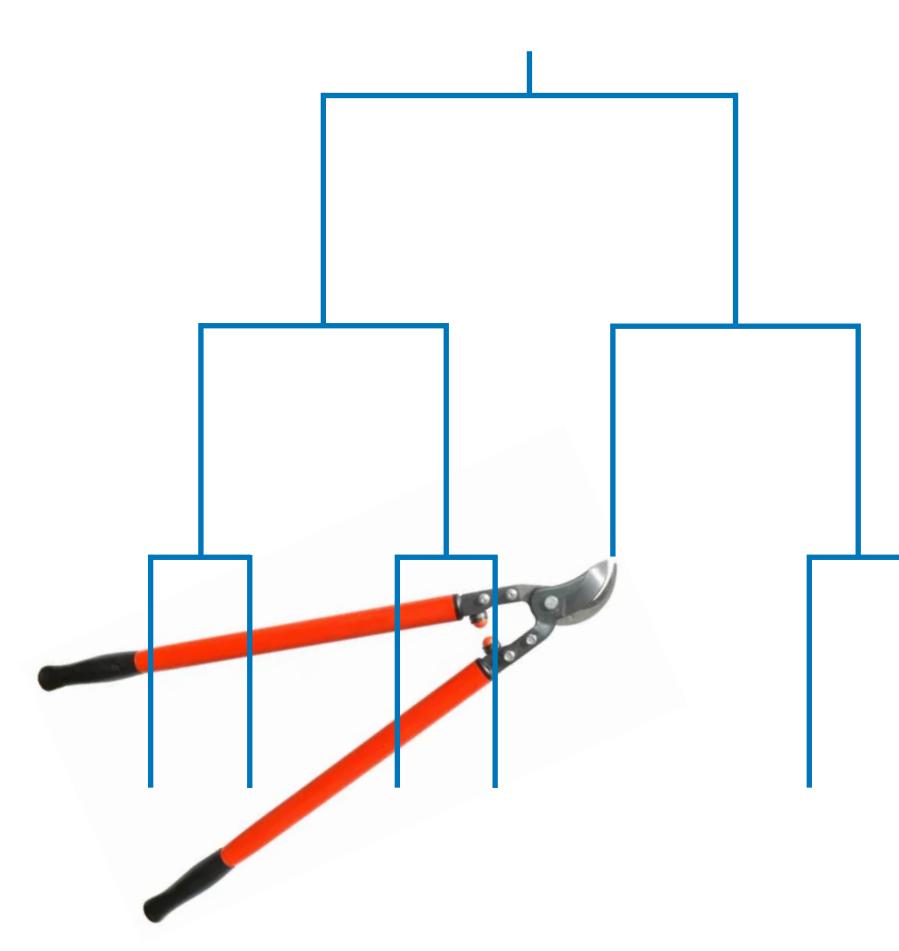
Idea: Recursively prune "weakest link" splits.





Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

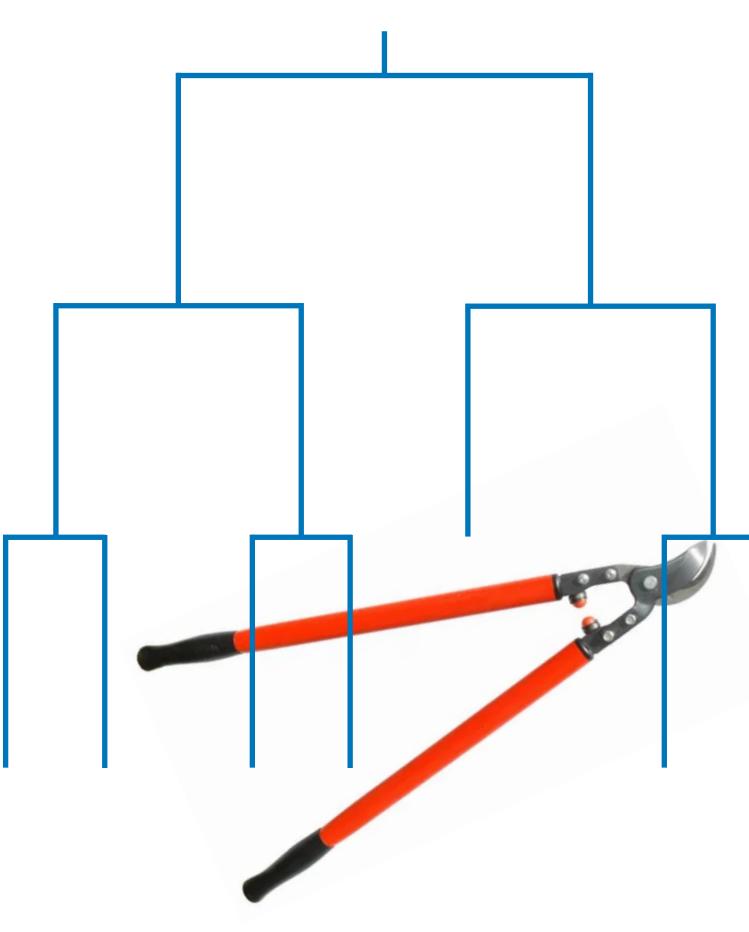
Idea: Recursively prune "weakest link" splits.





Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

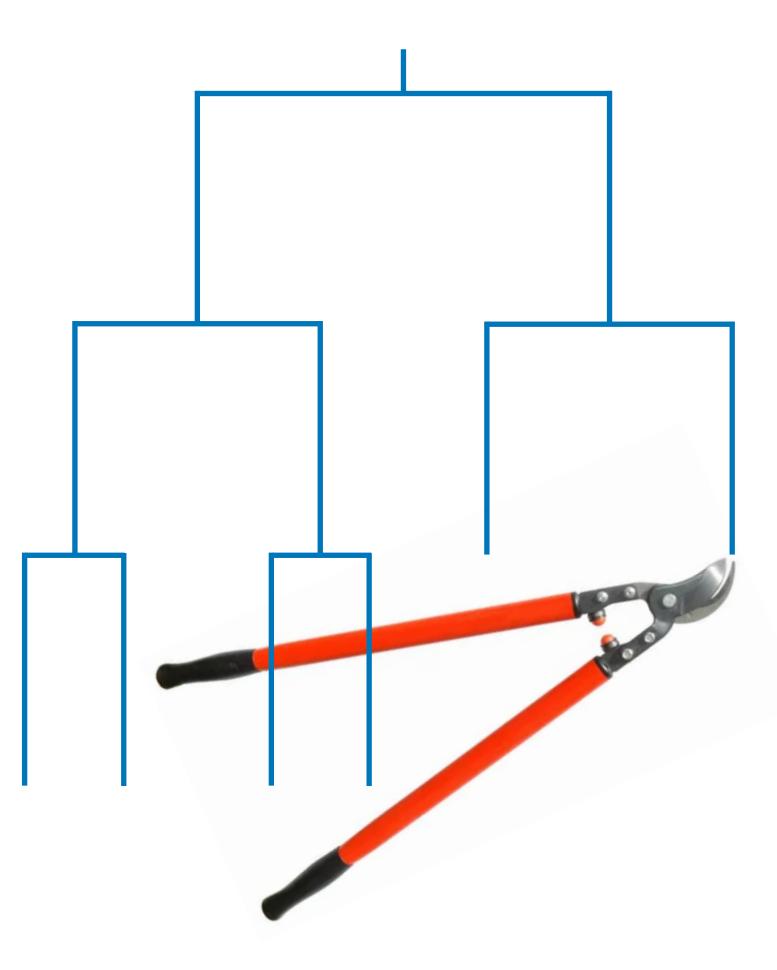
Idea: Recursively prune "weakest link" splits.





Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

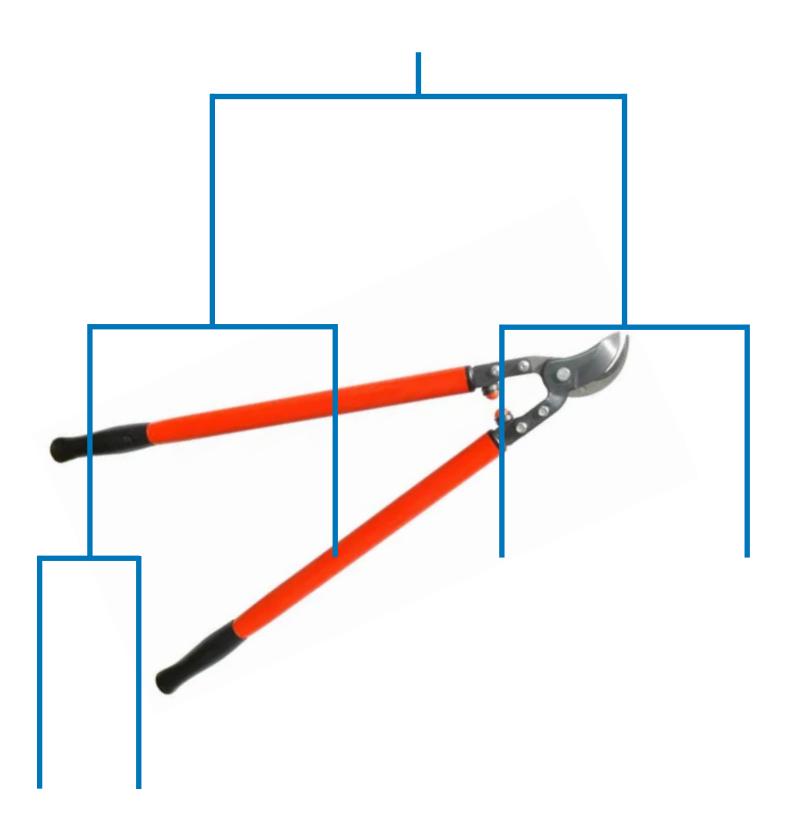
Idea: Recursively prune "weakest link" splits.

Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.

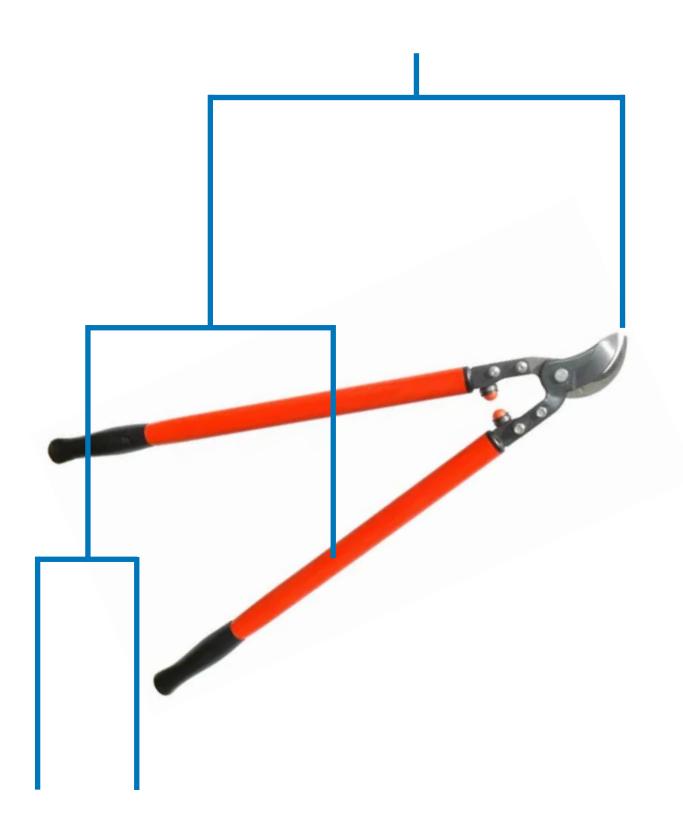
Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

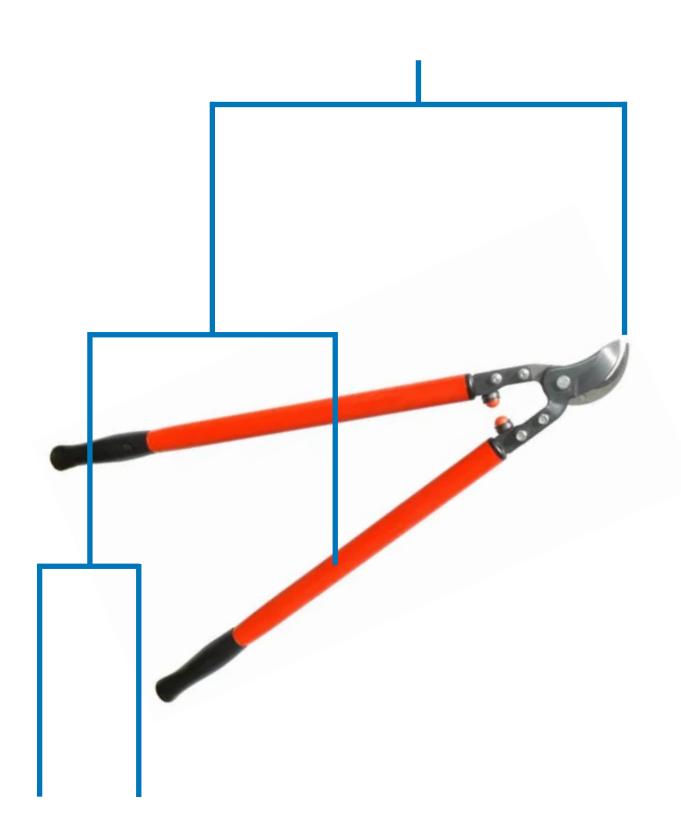
Idea: Recursively prune "weakest link" splits.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.

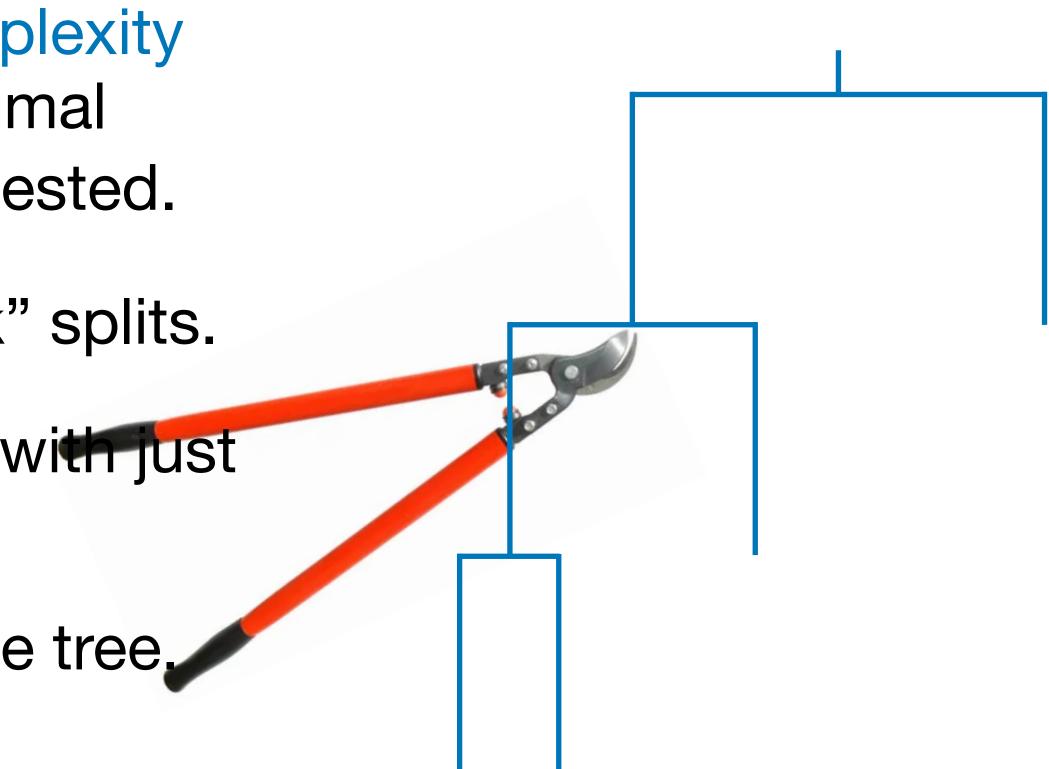
Usually, weakest link splits are those with just two terminal nodes below them.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.

Usually, weakest link splits are those with just two terminal nodes below them.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested. Idea: Recursively prune "weakest link" splits. Usually, weakest link splits are those with just two terminal nodes below them. Sometimes, they can be further up the tree.

Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.

Usually, weakest link splits are those with just two terminal nodes below them.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.

Usually, weakest link splits are those with just two terminal nodes below them.



Given a fully grown tree  $T_0$ , cost complexity pruning is an algorithm that finds optimal sequence  $T_{\alpha}$ , which turns out to be nested.

Idea: Recursively prune "weakest link" splits.

Usually, weakest link splits are those with just two terminal nodes below them.

• Grow a full tree  $T_0$  on the whole training data

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into *K* folds

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into *K* folds
- For each fold k,

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into *K* folds
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into *K* folds
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into K folds
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )
  - using these trees, make predictions for each in-fold observation i and each  $\alpha$

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into K folds
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )
  - using these trees, make predictions for each in-fold observation i and each  $\alpha$
- Find CV estimates and standard errors as usual; choose  $\hat{\alpha}$  based on 1-standard-error rule

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into K folds
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )
  - using these trees, make predictions for each in-fold observation i and each  $\alpha$
- Find CV estimates and standard errors as usual; choose  $\hat{\alpha}$  based on 1-standard-error rule
- Output the final decision tree  $T_{\hat{\alpha}}$  (based on sequence of trees grown on full training data)

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$
- Split the training samples into K folds
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )
  - using these trees, make predictions for each in-fold observation i and each  $\alpha$
- Find CV estimates and standard errors as usual; choose  $\hat{\alpha}$  based on 1-standard-error rule
- Output the final decision tree  $T_{\hat{\alpha}}$  (based on sequence of trees grown on full training data)

• Key insight: Cross-validating to find optimal  $\alpha$ ; trees with same  $\alpha$  across CV folds may be different.

- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$ • Analogy with lasso: The variables selected for the same  $\lambda$  across • Split the training samples into K folds different CV folds might be different.
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )
  - using these trees, make predictions for each in-fold observation i and each  $\alpha$
- Find CV estimates and standard errors as usual; choose  $\hat{\alpha}$  based on 1-standard-error rule
- Output the final decision tree  $T_{\hat{\alpha}}$  (based on sequence of trees grown on full training data)

• Key insight: Cross-validating to find optimal  $\alpha$ ; trees with same  $\alpha$  across CV folds may be different.

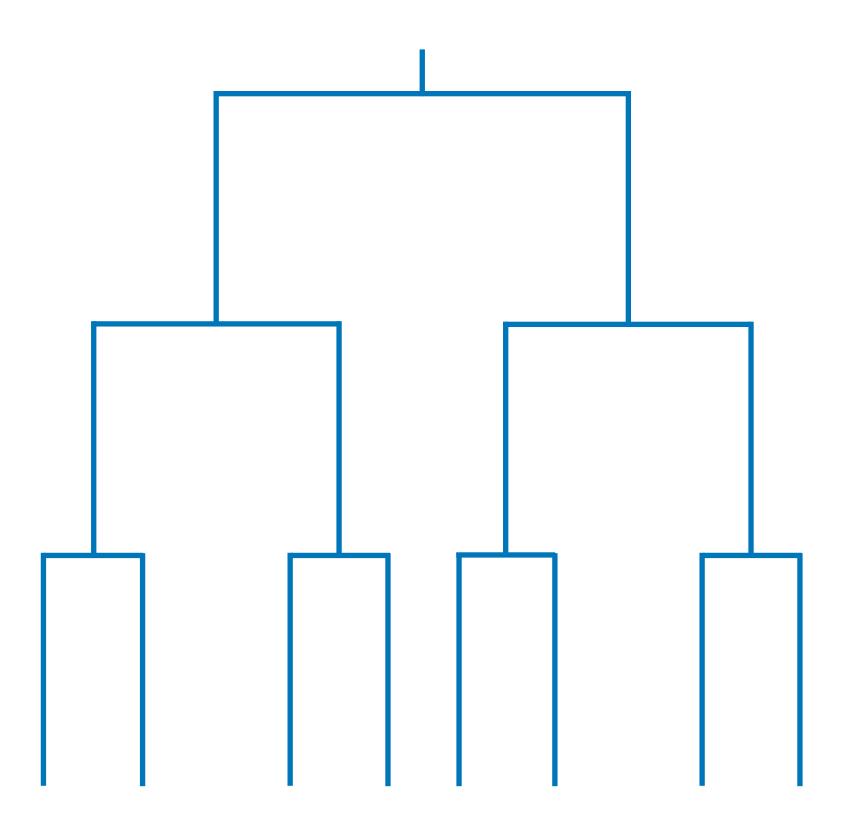
- Grow a full tree  $T_0$  on the whole training data
- Prune the tree to get a discrete sequence of trees  $T_{\alpha}$ • Analogy with lasso: The variables selected for the same  $\lambda$  across • Split the training samples into K folds different CV folds might be different.
- For each fold k,
  - grow a full tree  $T_0^{-k}$  on the out-of-fold data
  - find the sequence of subtrees  $T_{\alpha}^{-k}$  of  $T_{0}^{-k}$  by pruning (but using same  $\alpha$ )
  - using these trees, make predictions for each in-fold observation i and each  $\alpha$
- Find CV estimates and standard errors as usual; choose  $\hat{\alpha}$  based on 1-standard-error rule
- Output the final decision tree  $T_{\hat{\alpha}}$  (based on sequence of trees grown on full training data)

 Key insight: Cross-validating to find optimal  $\alpha$ ; trees with same  $\alpha$  across CV folds may be different.

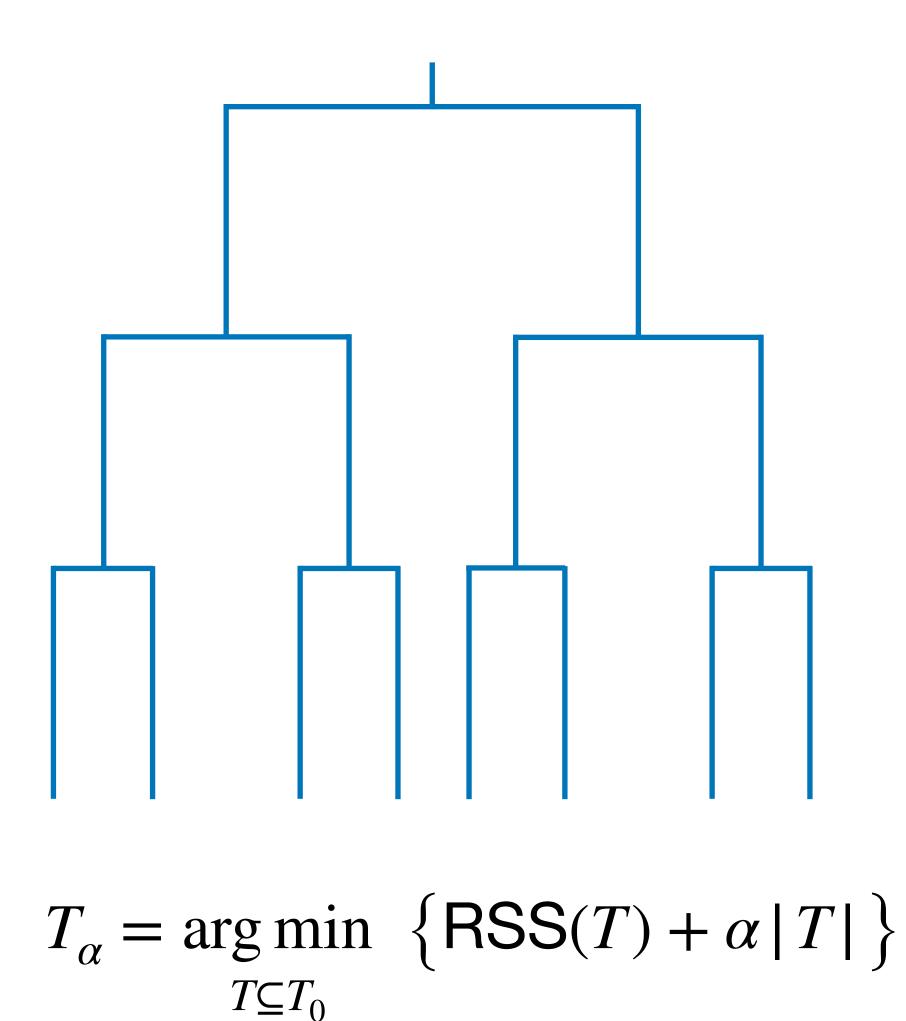
• Fitting to entire training data happens at the beginning of the process (for trees) rather than at the end (for lasso).



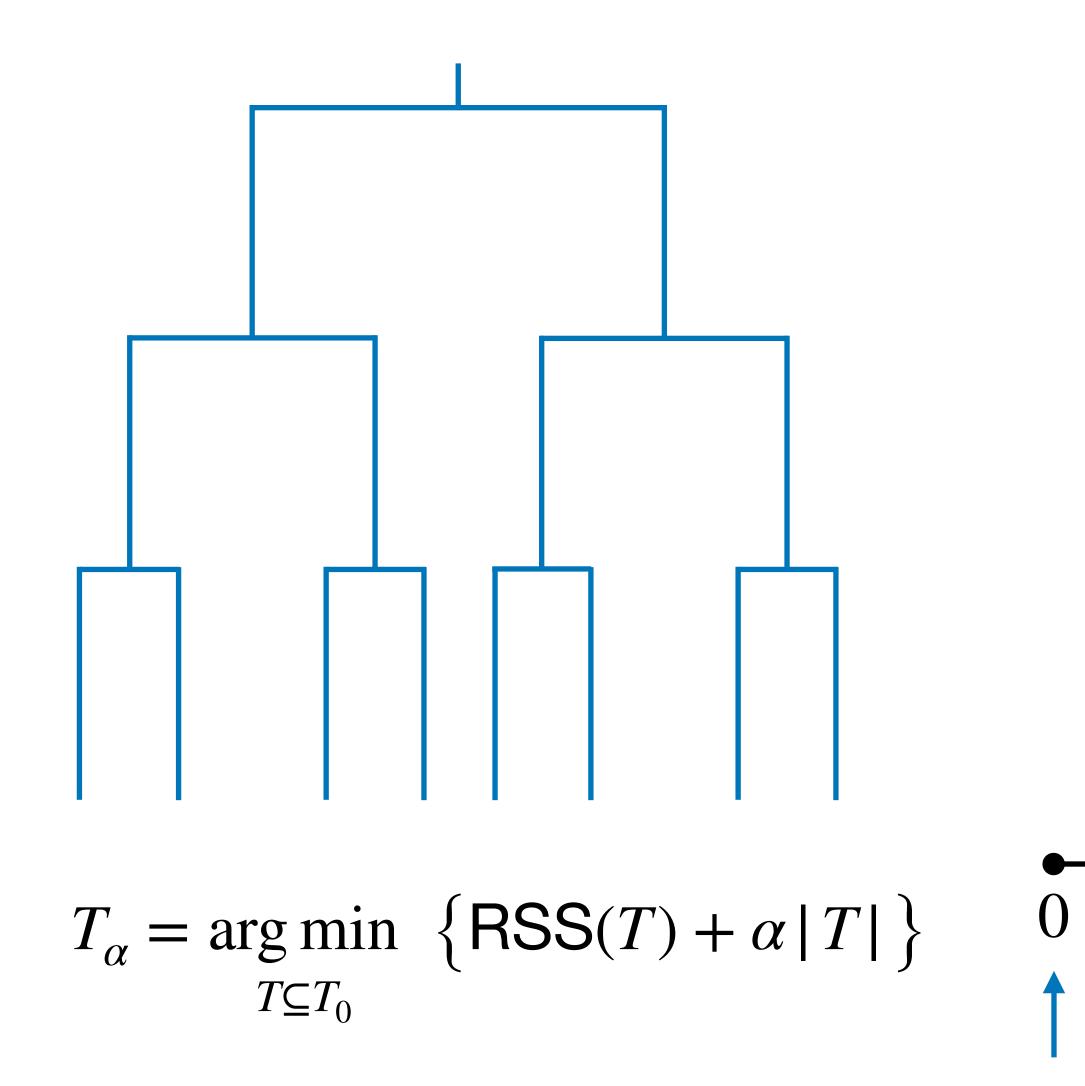
Step 1: Grow tree on whole training data in greedy fashion to get  $T_0$ .



## **Illustration: Tree growing and pruning** Step 2: Consider penalized objective function.



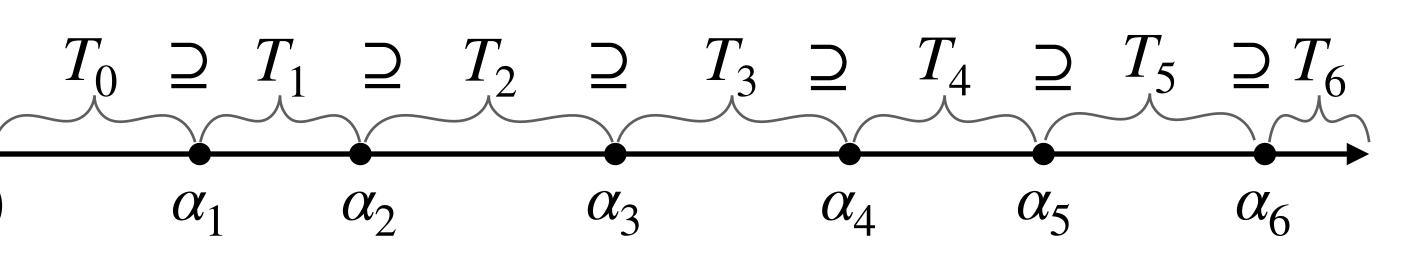
Step 3: Sweep  $\alpha$  from 0 to infinity, giving a nested sequence of trees.





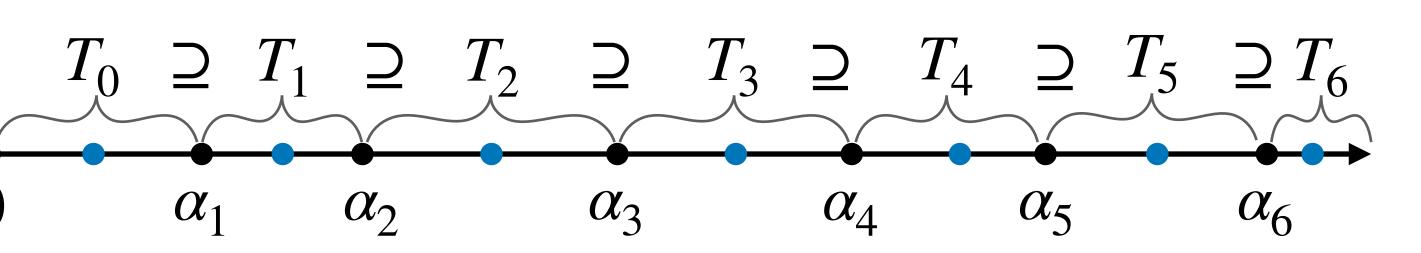
Step 3: Sweep  $\alpha$  from 0 to infinity, giving a nested sequence of trees.

 $T_{\alpha} = \underset{T \subseteq T_{0}}{\operatorname{arg\,min}} \left\{ \operatorname{RSS}(T) + \alpha | T | \right\} \qquad 0$ 



Step 4: Choose a representative value of  $\alpha$  for each tree.

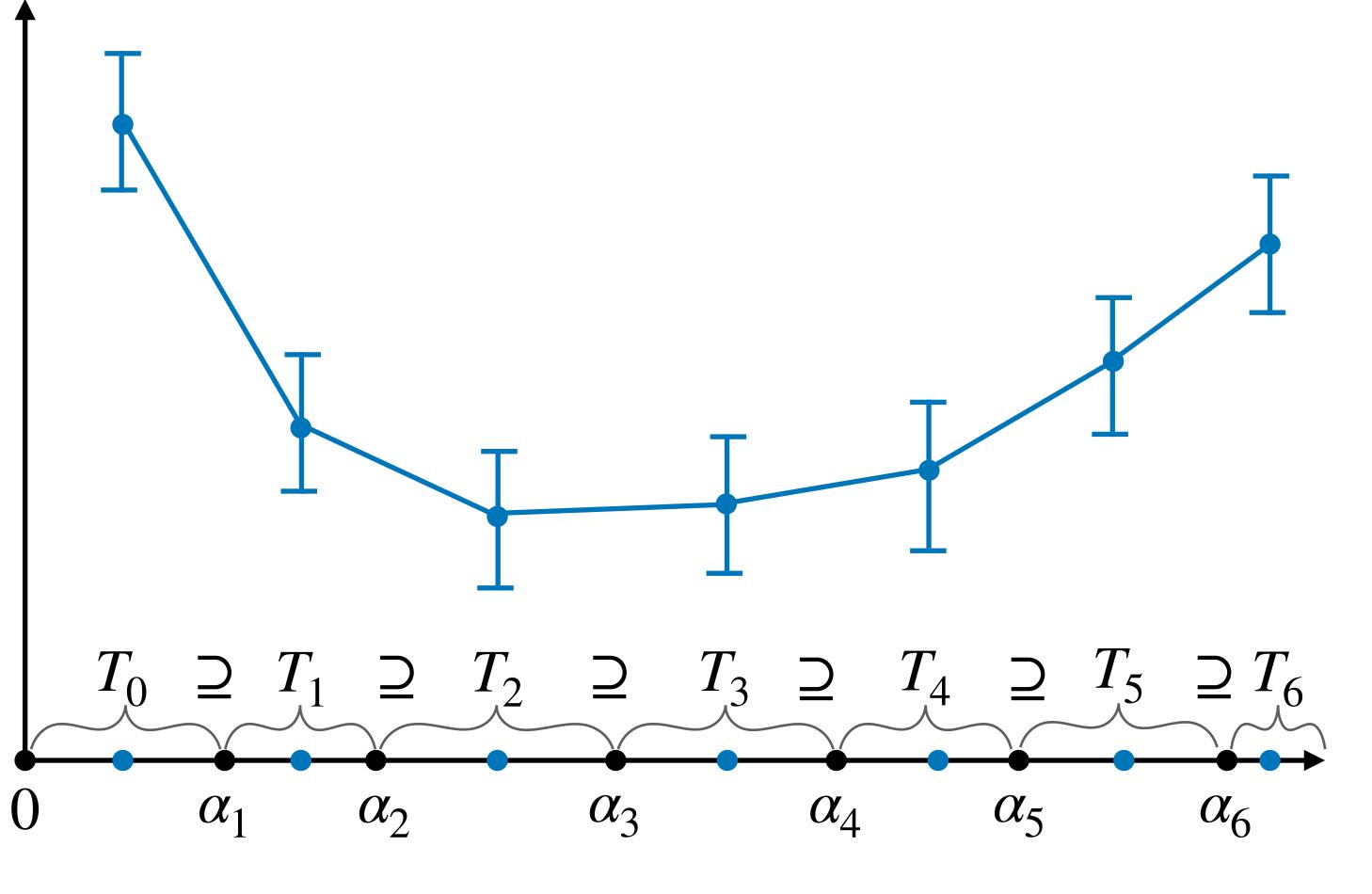
 $T_{\alpha} = \underset{T \subseteq T_{0}}{\operatorname{arg\,min}} \left\{ \operatorname{RSS}(T) + \alpha |T| \right\} \qquad 0$ 



Step 5: Cross-validate over the representative values of  $\alpha$ .

**CV error** 

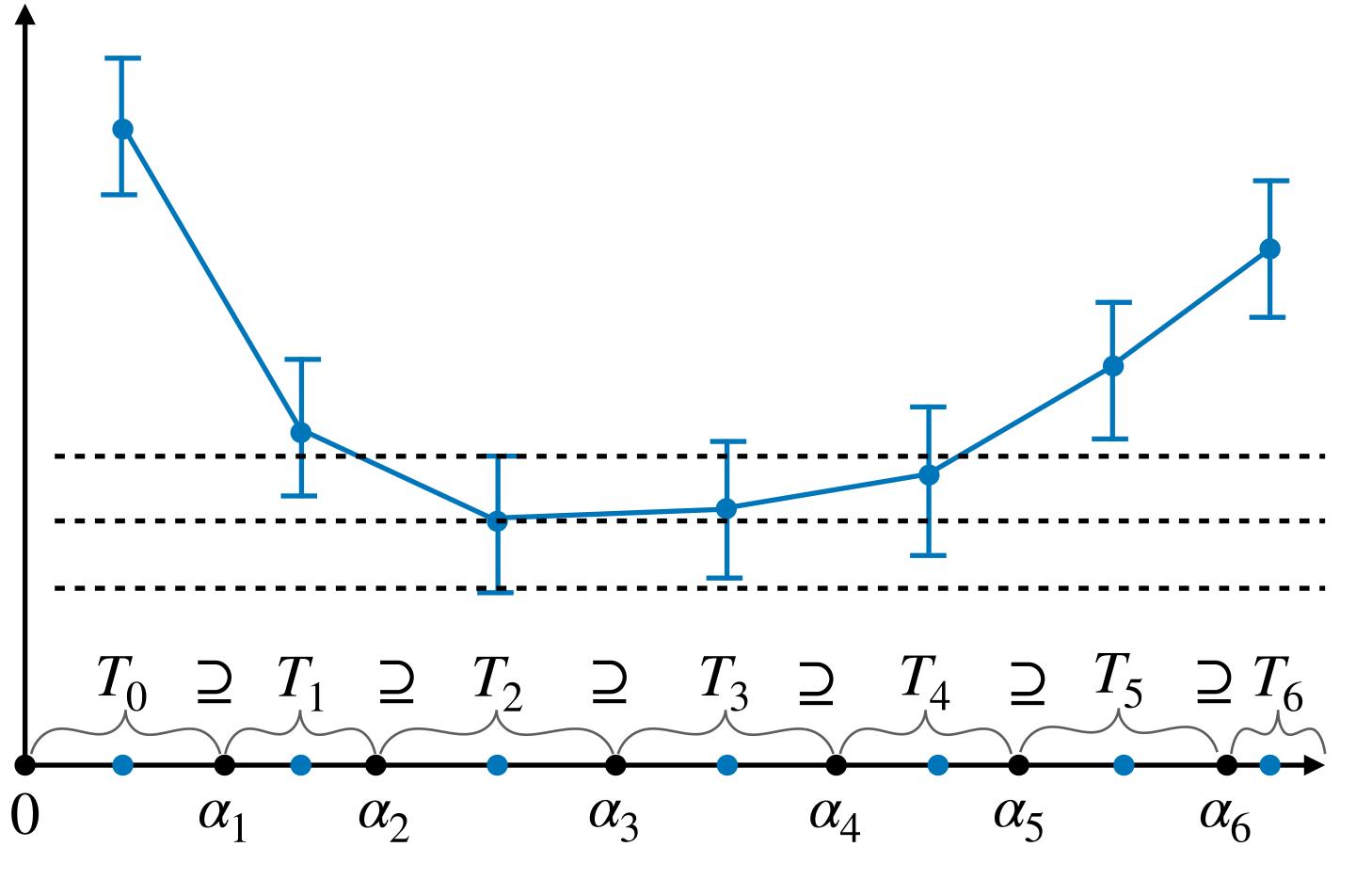
 $T_{\alpha} = \underset{T \subseteq T_{0}}{\operatorname{arg\,min}} \left\{ \operatorname{RSS}(T) + \alpha | T | \right\}$ 



Step 6: Use one-standard-error rule to choose  $\alpha$ .

error

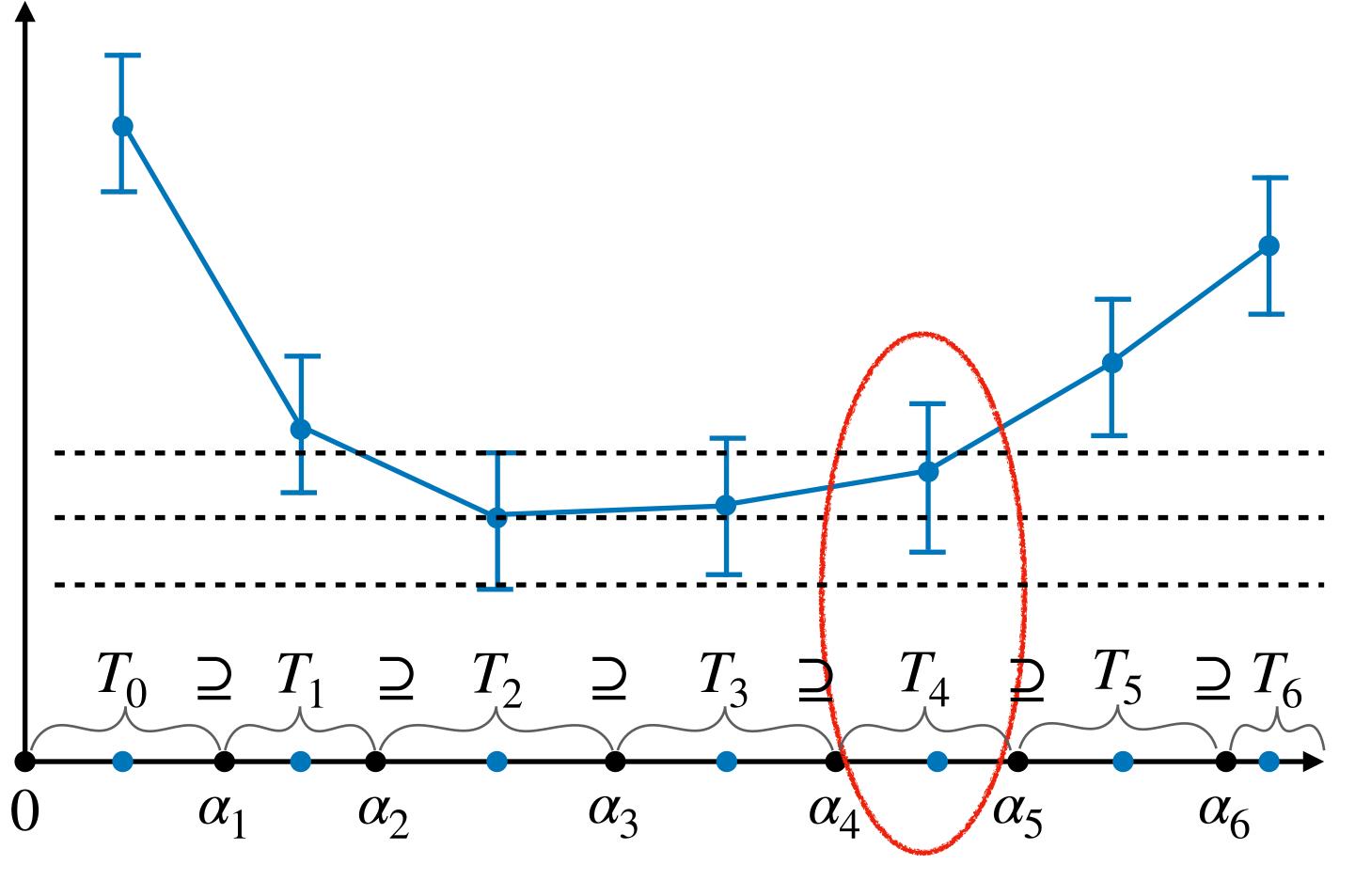
 $T_{\alpha} = \arg\min \left\{ \mathsf{RSS}(T) + \alpha |T| \right\}$  $T \subseteq T_0$ 



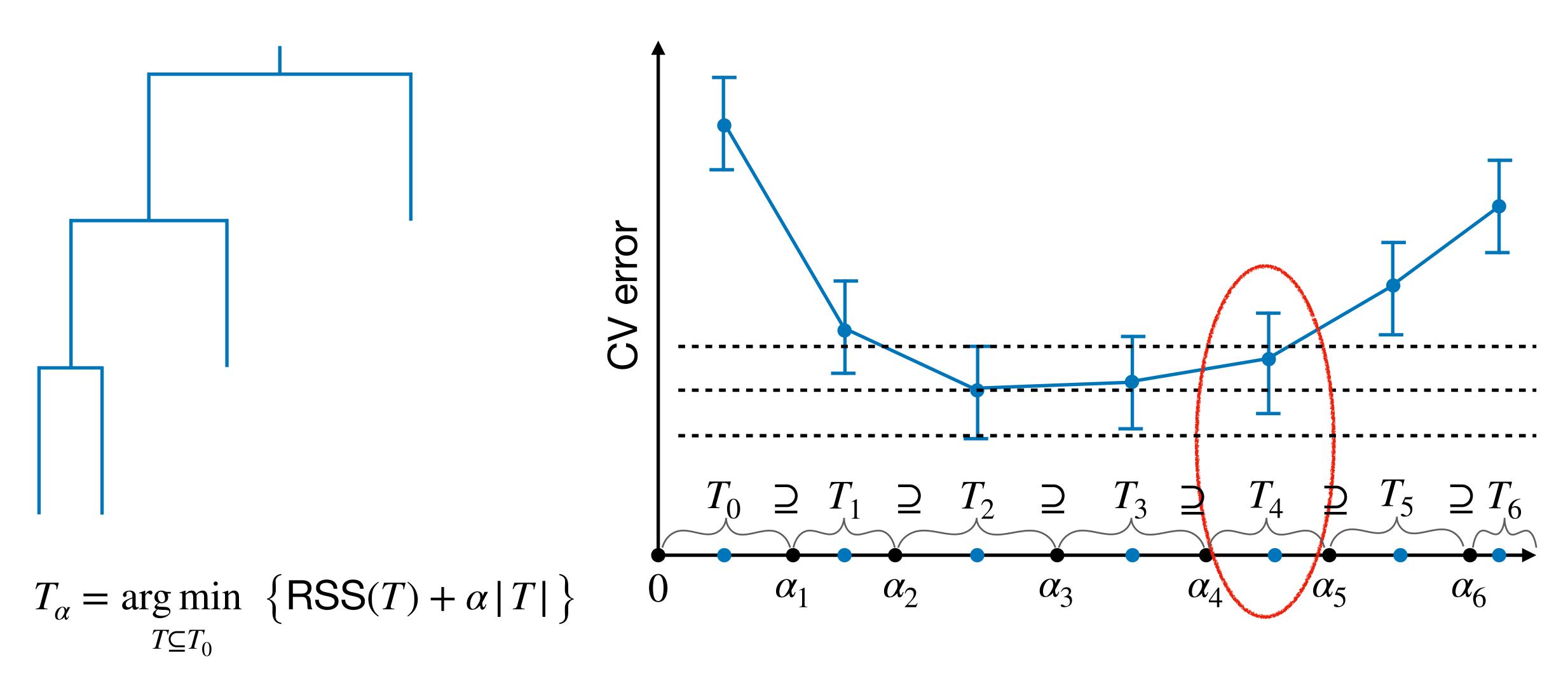
Step 6: Use one-standard-error rule to choose  $\alpha$ .

error

 $T_{\alpha} = \arg\min \left\{ \mathsf{RSS}(T) + \alpha |T| \right\}$  $T \subseteq T_0$ 



Step 6: Use one-standard-error rule to choose  $\alpha$ .



### Growing proceeds from smaller to larger trees; pruning from larger to smaller.

For regression trees, growing and pruning are both based on RSS.

Growing proceeds from smaller to larger trees; pruning from larger to smaller.

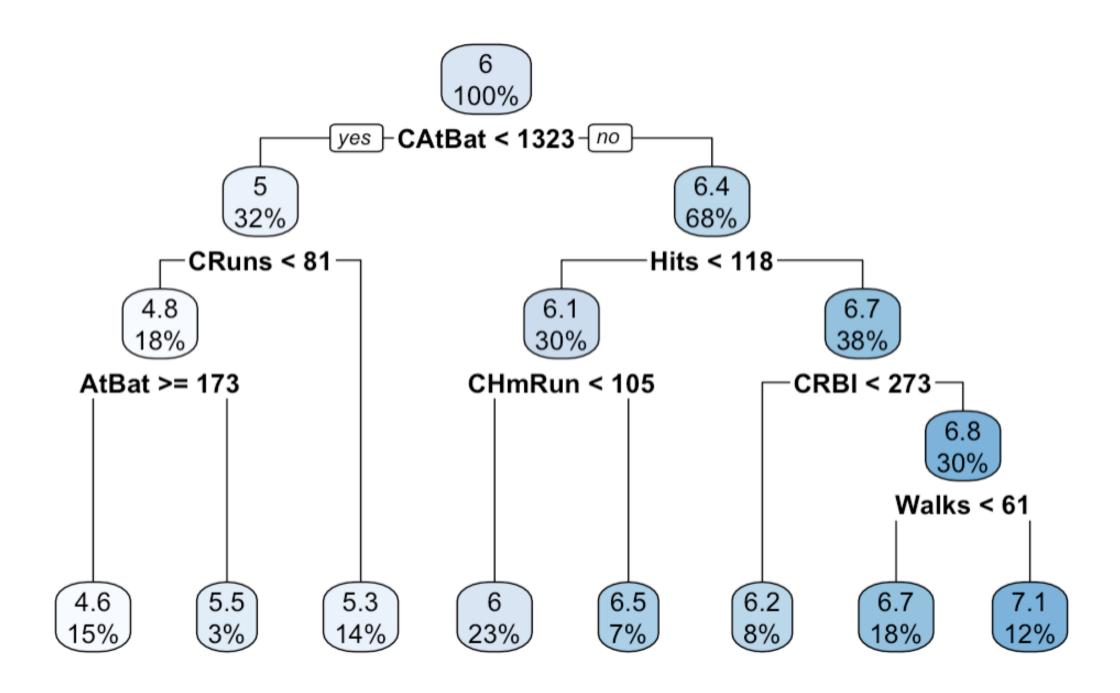
For regression trees, growing and pruning are both based on RSS.

For classification trees, growing based on Gini index but pruning based on misclassification error.

- Growing proceeds from smaller to larger trees; pruning from larger to smaller.

- Growing proceeds from smaller to larger trees; pruning from larger to smaller.
- For regression trees, growing and pruning are both based on RSS.
- For classification trees, growing based on Gini index but pruning based on misclassification error.
- Growing and pruning both define a sequence of trees, but it may not be the same sequence. The sequence of trees for growing not used except to get the big tree  $T_0$ ; the sequence of trees for pruning is the one used for cross-validation.

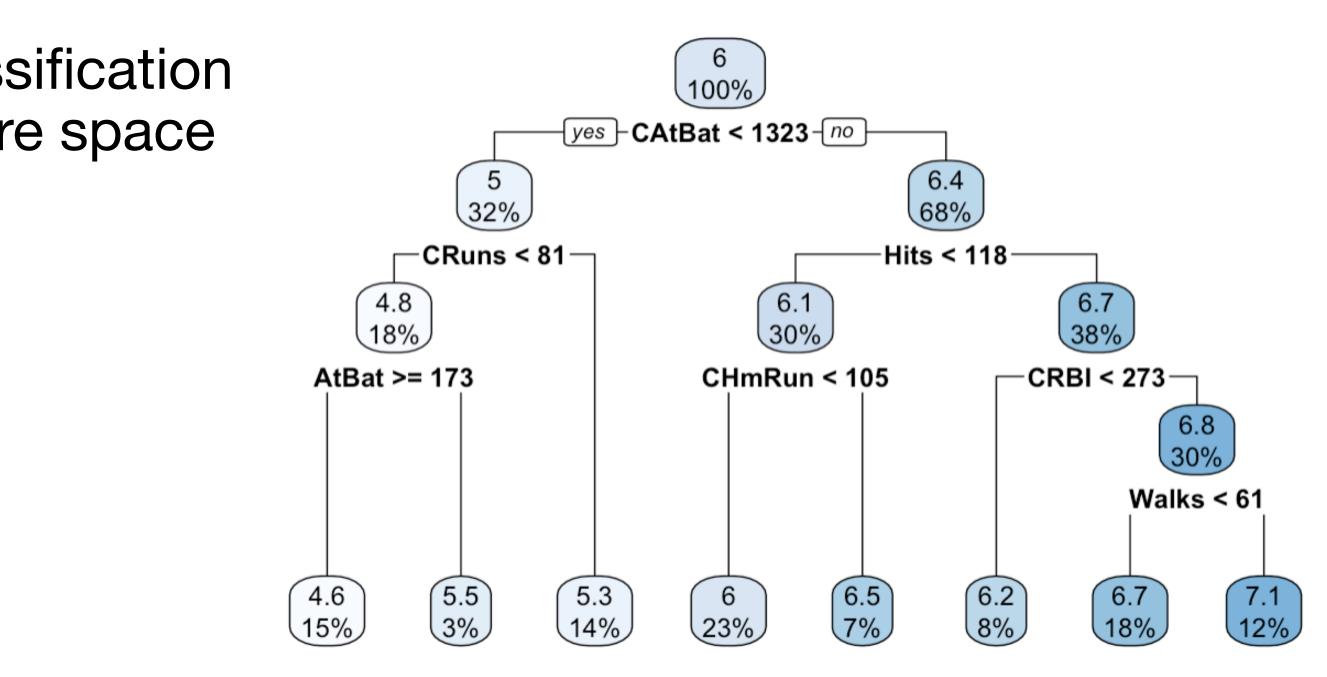
## **Summary: Decision Trees**



## **Summary: Decision Trees**

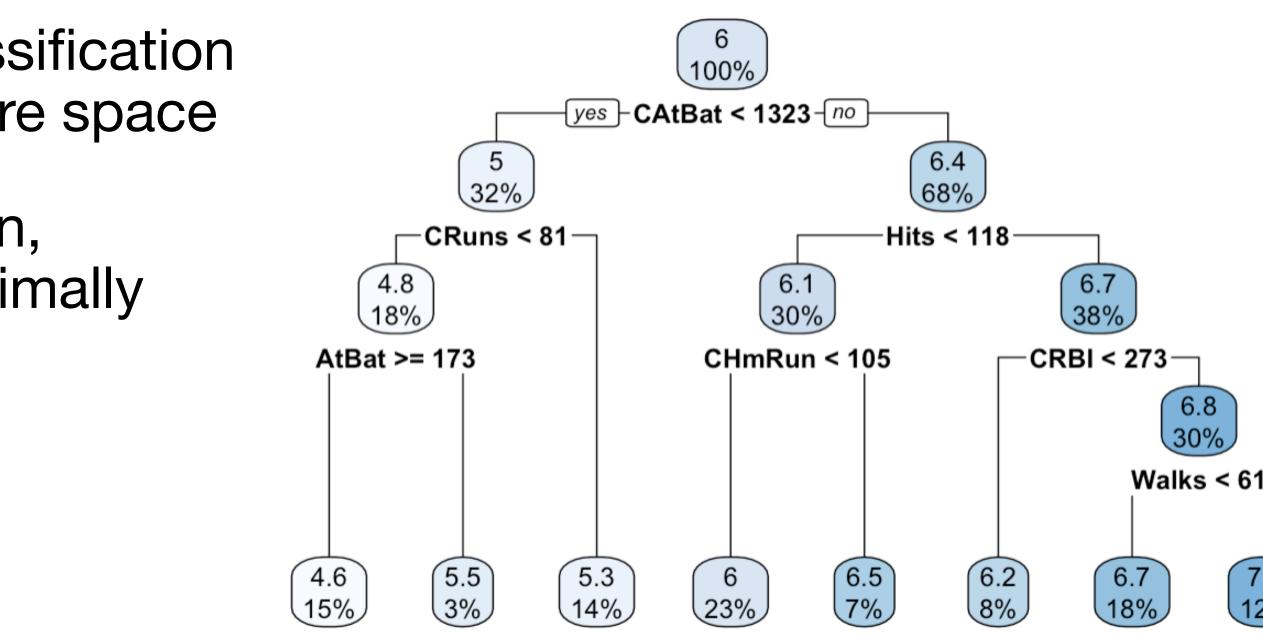
 Nonlinear method for regression or classification based on recursive partitioning of feature space

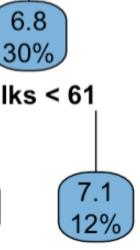




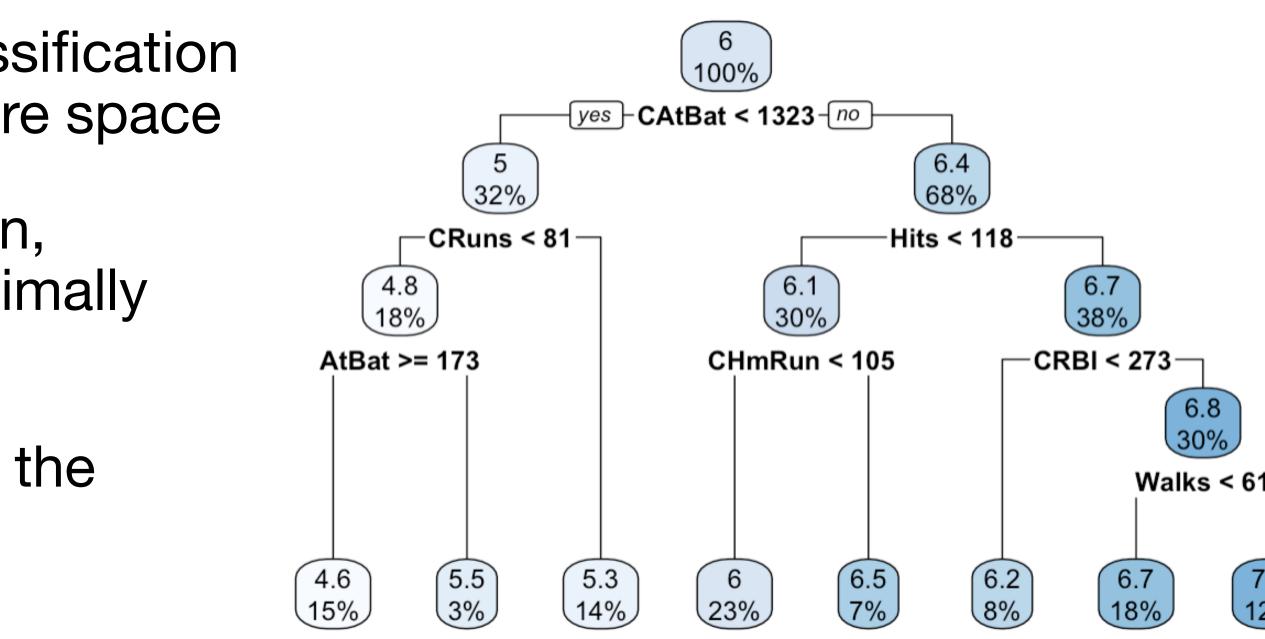
# **Summary: Decision Trees**

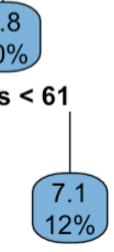
- Nonlinear method for regression or classification based on recursive partitioning of feature space
- Trees grown in greedy top-down fashion, choosing feature and split point to maximally improve purity of terminal nodes.



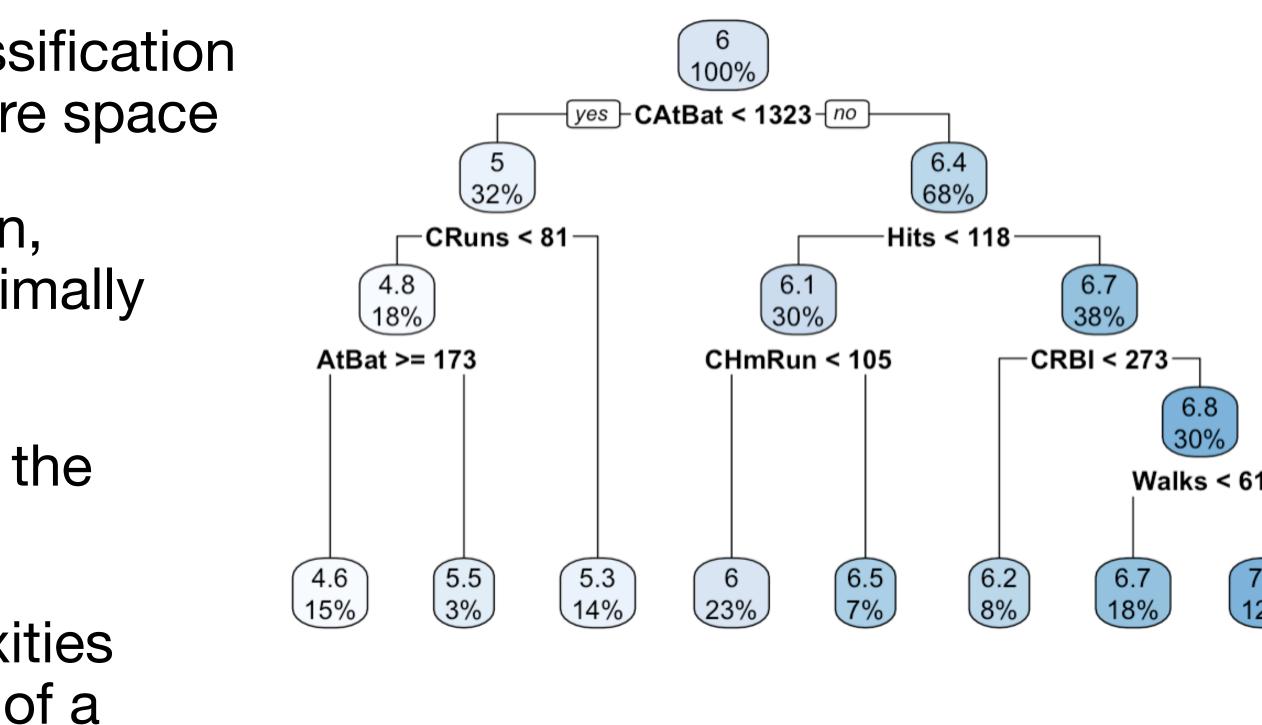


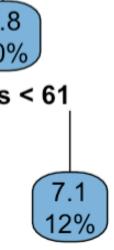
- Nonlinear method for regression or classification based on recursive partitioning of feature space
- Trees grown in greedy top-down fashion, choosing feature and split point to maximally improve purity of terminal nodes.
- The complexity of a tree increases with the number of terminal nodes.



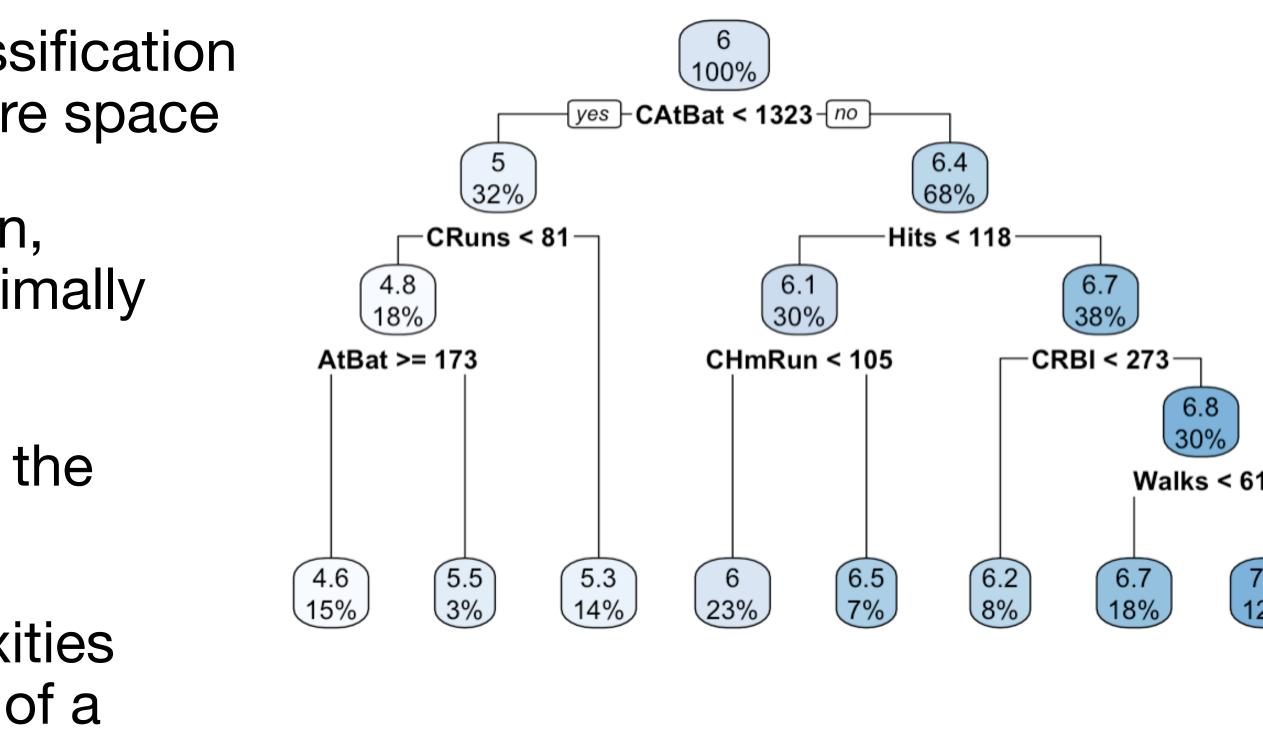


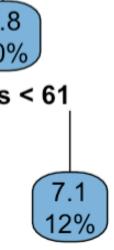
- Nonlinear method for regression or classification based on recursive partitioning of feature space
- Trees grown in greedy top-down fashion, choosing feature and split point to maximally improve purity of terminal nodes.
- The complexity of a tree increases with the number of terminal nodes.
- A sequence of trees of varying complexities obtained from cost complexity pruning of a maximally-grown tree.



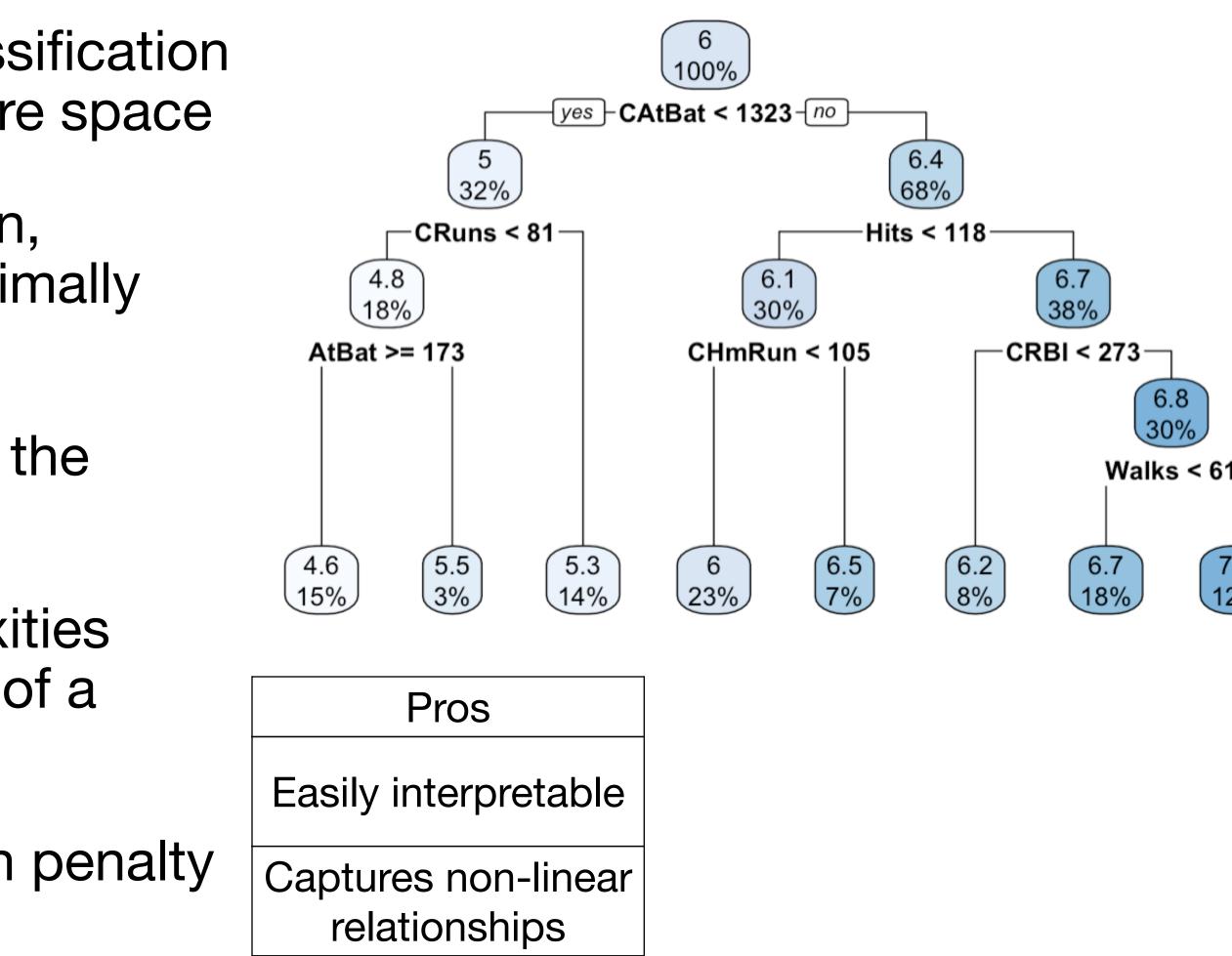


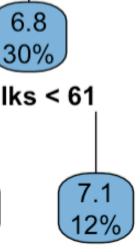
- Nonlinear method for regression or classification based on recursive partitioning of feature space
- Trees grown in greedy top-down fashion, choosing feature and split point to maximally improve purity of terminal nodes.
- The complexity of a tree increases with the number of terminal nodes.
- A sequence of trees of varying complexities obtained from cost complexity pruning of a maximally-grown tree.
- Final tree chosen by cross-validation on penalty parameter  $\alpha$ .



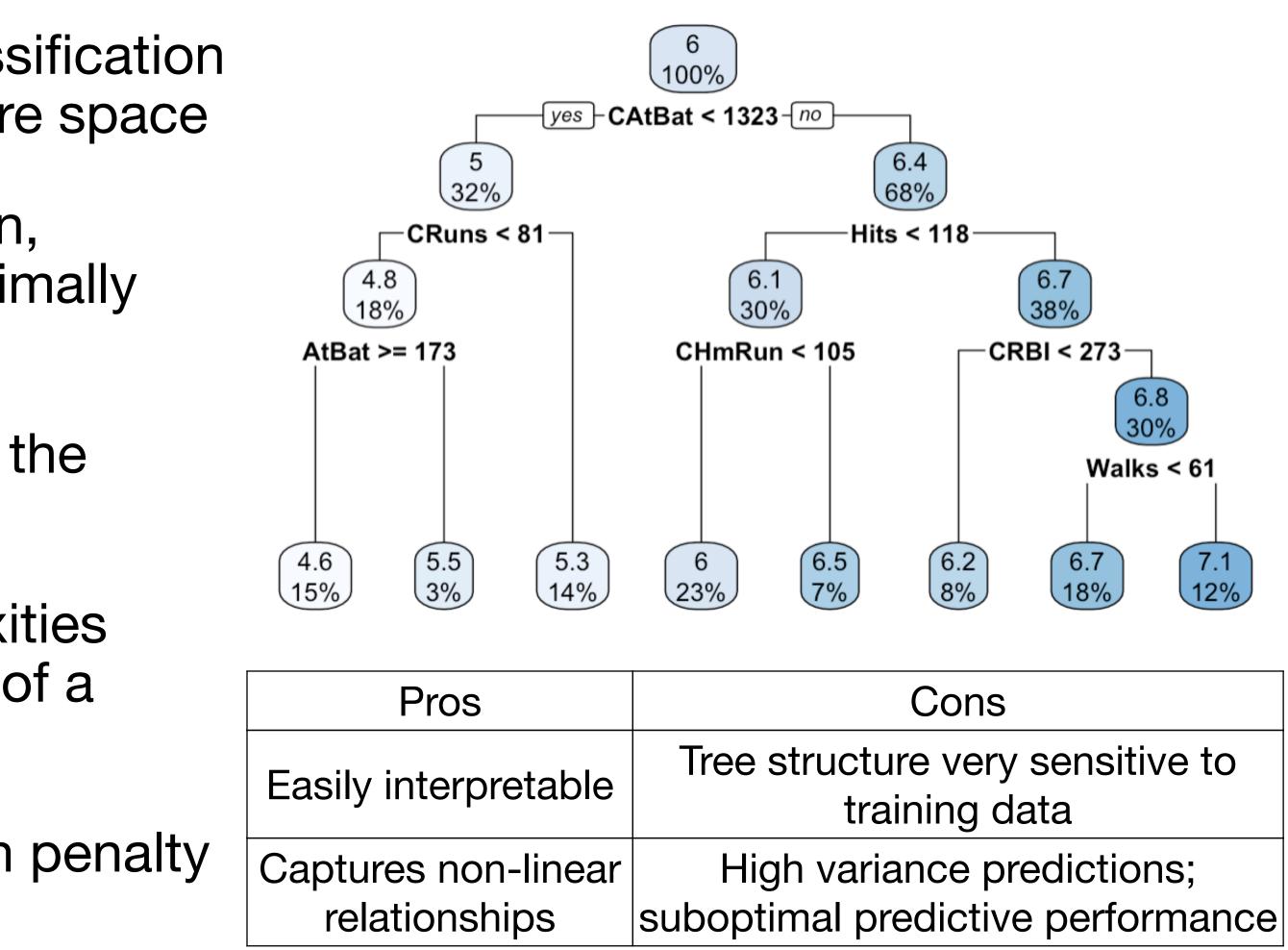


- Nonlinear method for regression or classification based on recursive partitioning of feature space
- Trees grown in greedy top-down fashion, choosing feature and split point to maximally improve purity of terminal nodes.
- The complexity of a tree increases with the number of terminal nodes.
- A sequence of trees of varying complexities obtained from cost complexity pruning of a maximally-grown tree.
- Final tree chosen by cross-validation on penalty parameter  $\alpha$ .





- Nonlinear method for regression or classification based on recursive partitioning of feature space
- Trees grown in greedy top-down fashion, choosing feature and split point to maximally improve purity of terminal nodes.
- The complexity of a tree increases with the number of terminal nodes.
- A sequence of trees of varying complexities obtained from cost complexity pruning of a maximally-grown tree.
- Final tree chosen by cross-validation on penalty parameter  $\alpha$ .



When it comes to prediction accuracy, trees suffer because of their high variance.



- When it comes to prediction accuracy, trees suffer because of their high variance.
- Here's an idea for how we can obtain a prediction method with lower variance:



• "Shake up" the training data lots of times (bootstrap)

- When it comes to prediction accuracy, trees suffer because of their high variance.
- Here's an idea for how we can obtain a prediction method with lower variance:



- "Shake up" the training data lots of times (bootstrap)
- For each version of the training data, fit a different tree

- When it comes to prediction accuracy, trees suffer because of their high variance.
- Here's an idea for how we can obtain a prediction method with lower variance:



- "Shake up" the training data lots of times (bootstrap)
- For each version of the training data, fit a different tree
- Use the average of all these trees to make predictions (aggregation)

- When it comes to prediction accuracy, trees suffer because of their high variance.
- Here's an idea for how we can obtain a prediction method with lower variance:



- "Shake up" the training data lots of times (bootstrap)
- For each version of the training data, fit a different tree
- Use the average of all these trees to make predictions (aggregation)
  - Bagging = Bootstrap Aggregation.

- When it comes to prediction accuracy, trees suffer because of their high variance.
- Here's an idea for how we can obtain a prediction method with lower variance:



- "Shake up" the training data lots of times (bootstrap)
- For each version of the training data, fit a different tree
- Use the average of all these trees to make predictions (aggregation)
  - Bagging = Bootstrap Aggregation.

- When it comes to prediction accuracy, trees suffer because of their high variance.
- Here's an idea for how we can obtain a prediction method with lower variance:

Intuition: By averaging a bunch of trees, we are reducing the variance while keeping the bias about the same. This should yield better predictive performance!







What we ideally would have wanted is to get many different random realizations of the training data, on which we could fit different trees.

of the training data, on which we could fit different trees.

random versions of our data by bootstrapping:

- What we ideally would have wanted is to get many different random realizations
- We only get one realization of the training data, but we can still get different

of the training data, on which we could fit different trees.

random versions of our data by bootstrapping:

- What we ideally would have wanted is to get many different random realizations
- We only get one realization of the training data, but we can still get different
- A bootstrap sample is a new data set with the same number of observations, generating by sampling observations from the original data with replacement.

of the training data, on which we could fit different trees.

random versions of our data by bootstrapping:

- What we ideally would have wanted is to get many different random realizations
- We only get one realization of the training data, but we can still get different
- A bootstrap sample is a new data set with the same number of observations, generating by sampling observations from the original data with replacement.
- The idea is that your bootstrap samples are slightly different versions of your training data, allowing you to fit different trees to these different training sets.

### Original training data

<b>Observation ID</b>	X	Y
1	X1	<b>Y</b> <sub>1</sub>
2	X <sub>2</sub>	<b>Y</b> <sub>2</sub>
3	X3	<b>Y</b> 3
4	X4	Y <sub>4</sub>
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>

### Original training data

<b>Observation ID</b>	X	Y
1	X <sub>1</sub>	Y <sub>1</sub>
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	Y <sub>3</sub>
4	X4	Y4
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>



<b>Observation ID</b>	X	Y
-----------------------	---	---

### Original training data

<b>Observation ID</b>	X	Y
1	X1	Y <sub>1</sub>
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	Y <sub>3</sub>
4	X4	Y4
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>



<b>Observation ID</b>	X	Y
5	<b>X</b> 5	<b>Y</b> 5

### Original training data

<b>Observation ID</b>	X	Y
1	X1	Y <sub>1</sub>
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	Y <sub>3</sub>
4	X4	Y4
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>



<b>Observation ID</b>	X	Y
5	<b>X</b> 5	<b>Y</b> 5
3	<b>X</b> 3	<b>Y</b> 3

### Original training data

<b>Observation ID</b>	X	Y
1	X1	Y <sub>1</sub>
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	Y <sub>3</sub>
4	X4	Y4
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>



<b>Observation ID</b>	X	Y
5	X5	<b>Y</b> 5
3	X <sub>3</sub>	<b>Y</b> 3
2	X <sub>2</sub>	<b>Y</b> <sub>2</sub>

### Original training data

<b>Observation ID</b>	X	Y
1	X1	Y <sub>1</sub>
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	Y <sub>3</sub>
4	X4	Y4
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>

<b>Observation ID</b>	X	Y
5	X <sub>5</sub>	<b>Y</b> <sub>5</sub>
3	X <sub>3</sub>	<b>Y</b> <sub>3</sub>
2	X <sub>2</sub>	<b>Y</b> <sub>2</sub>
3	X3	<b>Y</b> <sub>3</sub>

### Original training data

<b>Observation ID</b>	X	Y
1	X <sub>1</sub>	Y <sub>1</sub>
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	Y <sub>3</sub>
4	X4	Y4
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>

<b>Observation ID</b>	X	Y
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>
3	X3	<b>Y</b> 3
2	X <sub>2</sub>	<b>Y</b> <sub>2</sub>
3	X3	<b>Y</b> 3
1	X <sub>1</sub>	<b>Y</b> <sub>1</sub>

### Original training data

<b>Observation ID</b>	X	Y	
1	X <sub>1</sub>	Y <sub>1</sub>	
2	X <sub>2</sub>	Y <sub>2</sub>	
3	X3	<b>Y</b> <sub>3</sub>	
4	X4	Y4	
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>	

#### Bootstrap sample 1

<b>Observation ID</b>	X	Y
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>
3	X <sub>3</sub>	<b>Y</b> 3
2	X <sub>2</sub>	Y <sub>2</sub>
3	X3	<b>Y</b> <sub>3</sub>
1	X <sub>1</sub>	<b>Y</b> <sub>1</sub>

#### Bootstrap sample *B*

<b>Observation ID</b>	X	Y
4	X4	Y4
1	X <sub>1</sub>	Y <sub>1</sub>
1	X <sub>1</sub>	Y <sub>1</sub>
5	<b>X</b> 5	<b>Y</b> <sub>5</sub>
4	X4	Y4

