

The background features a dense field of binary digits (0s and 1s) in a light blue/cyan color, creating a digital rain effect. Overlaid on this is a heatmap with a color gradient from dark blue to bright yellow, showing irregular, organic shapes that suggest spatial data or biological patterns.

# GG 501 SPATIAL KNOWLEDGE MOBILIZATION

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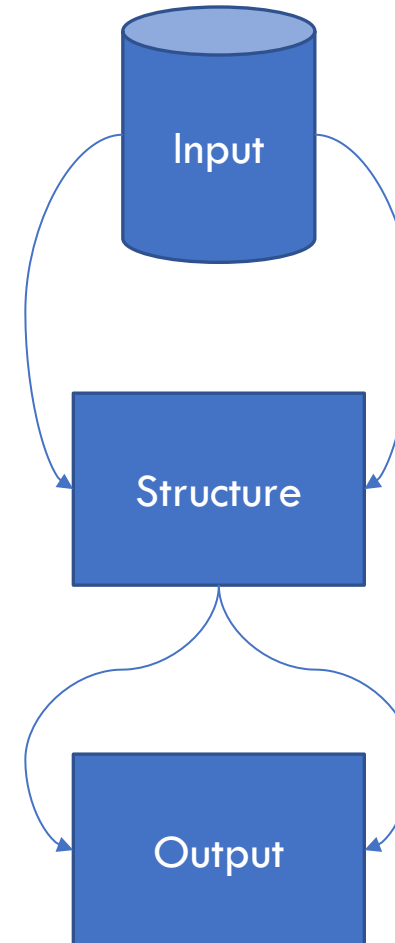
Mar 1: Parameterization and validation I

# MODELS — parameterization and validation

- Any time we fit a model, we have to make choices
  - What data goes in
  - What settings or configurations need to be set to run the model
    - this is parameterization
    - these are model-dependent
-

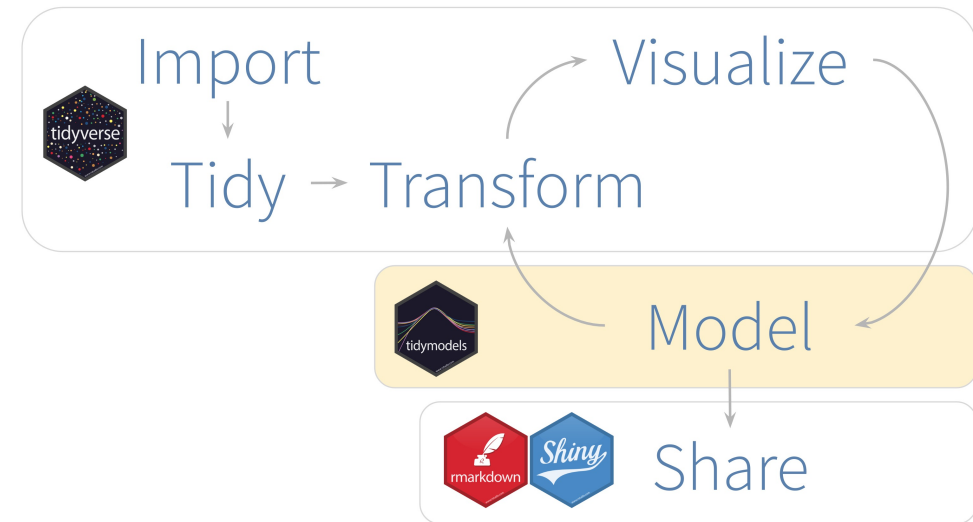
# MODELS COME IN MANY VARIETIES

- Each take some input data
- Attempt to generalize about the underlying data-generating-process
- Can be used for a variety of purposes –
  - description
  - explanation
  - prediction



# TIDYMODELS

- provide a clean and unified interface for modelling in data as part of an overall tidy workflow
- a collection of packages that focus on common aspects of statistical modelling and support many different versions of models implemented in different packages



Pre-Process → Train → Validate

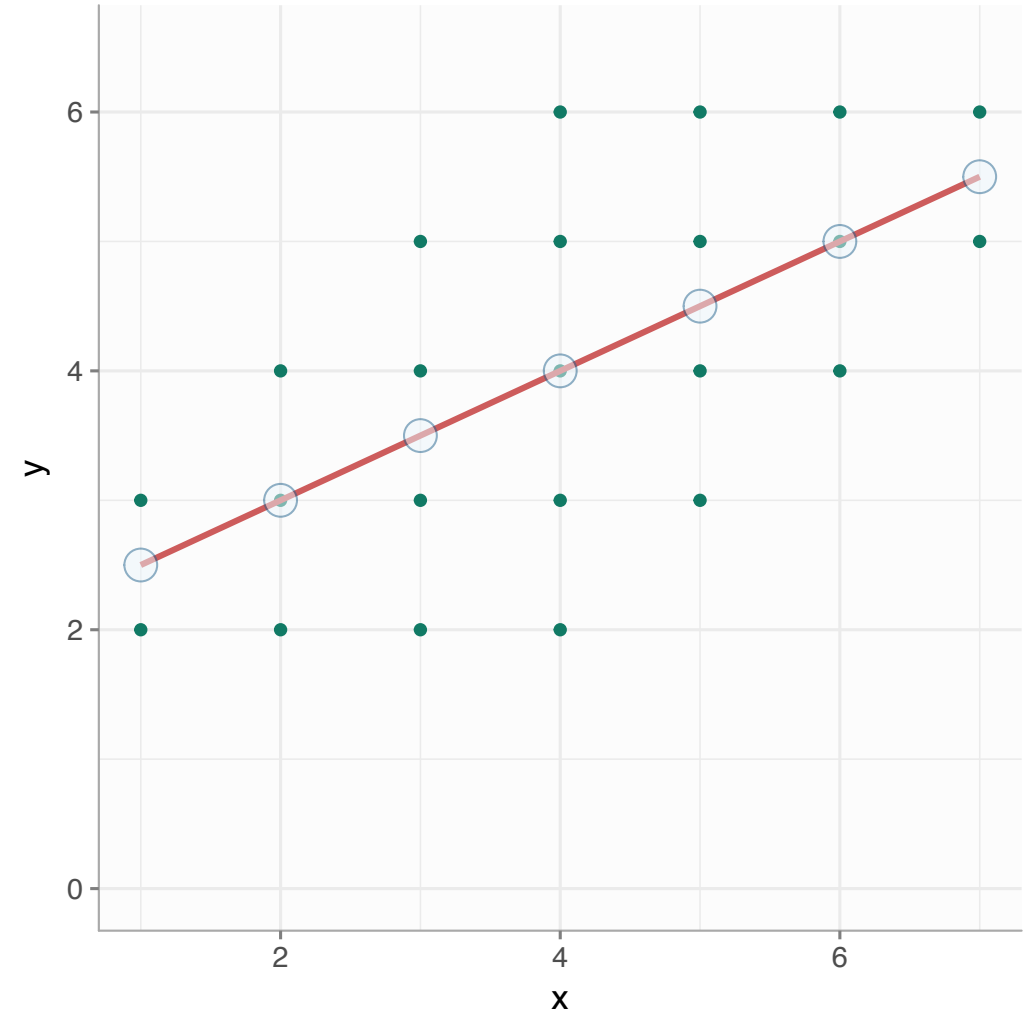


# LINEAR MODEL IN R `lm`

- simple linear regression model available in r function `lm`
- linear model fits a relationship between covariates and the conditional mean of the response or dependent variable
- has strict assumptions regarding independence of error terms which have implications for using with spatial / environmental data (and temporally correlated data)

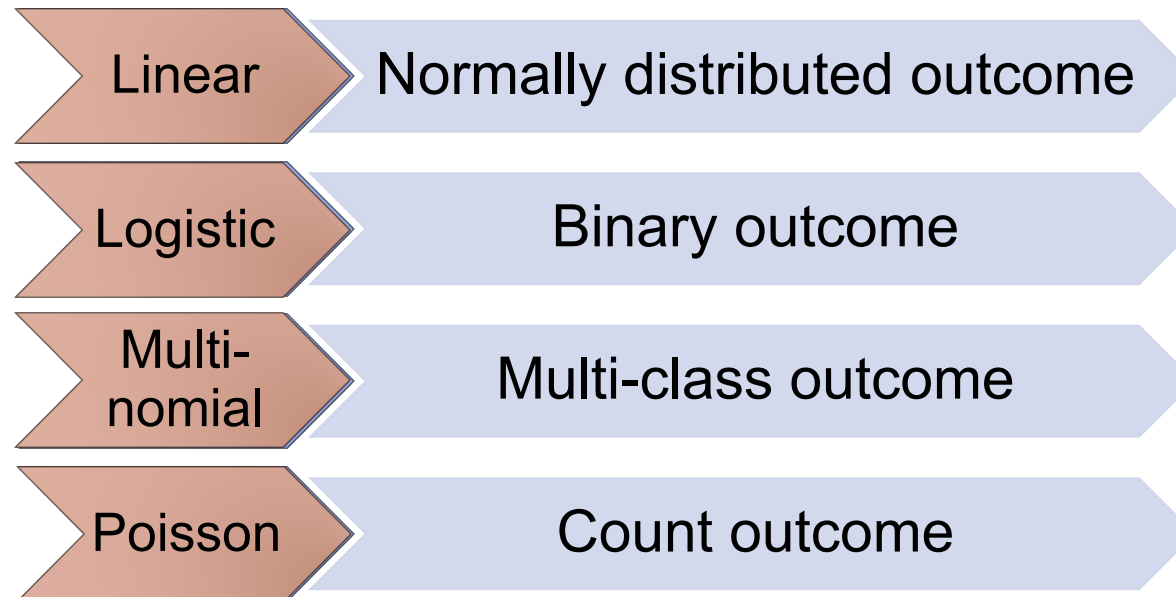
# LINEAR REGRESSION IN R

- Recalling `lm` in R
- Each point represents a single observation
- The red line is the **line of best fit**
  - all predicted values from this model will fall on the line of best fit
- The line goes through each *conditional mean*
  - It goes through the mean at each value of  $x$ 
    - E.g. When  $x = 1$ , mean of  $y = 2.5$  (the conditional mean of  $y$  at  $x = 1$  is 2.5)



# GENERALIZED LINEAR MODELS

- Flexible generalization of ordinary linear regression.
- Allows for outcomes that have other than a normal distribution.
- R implementation considers all models and link functions implemented in the R function `glm`



# glm IN R

```
## an example with offsets from Venables & Ripley (2002,
p.189)
utils::data(anorexia, package = "MASS")
anorex.1 <- glm(Postwt ~ Prewt + Treat + offset(Prewt),
family = gaussian, data = anorexia)
summary(anorex.1)
```

```
## Dobson (1990) Page 93: Randomized Controlled Trial :
counts <- c(18,17,15,20,10,20,25,13,12)
outcome <- gl(3,1,9)
treatment <- gl(3,3)
data.frame(treatment, outcome, counts)
# showing data
glm.D93 <- glm(counts ~ outcome + treatment, family =
poisson())
```

```
> summary(anorex.1)
```

Call:

```
glm(formula = Postwt ~ Prewt + Treat + offset(Prewt),
family = gaussian,
data = anorexia)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-14.1083	-4.2773	-0.5484	5.4838	15.2922

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	49.7711	13.3910	3.717	0.000410	***
Prewt	-0.5655	0.1612	-3.509	0.000803	***
TreatCont	-4.0971	1.8935	-2.164	0.033999	*
TreatFT	4.5631	2.1333	2.139	0.036035	*

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Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.'  
0.1 ' ' 1

(Dispersion parameter for gaussian family taken to be  
48.69504)

Null deviance: 4525.4 on 71 degrees of freedom  
Residual deviance: 3311.3 on 68 degrees of freedom  
AIC: 489.97

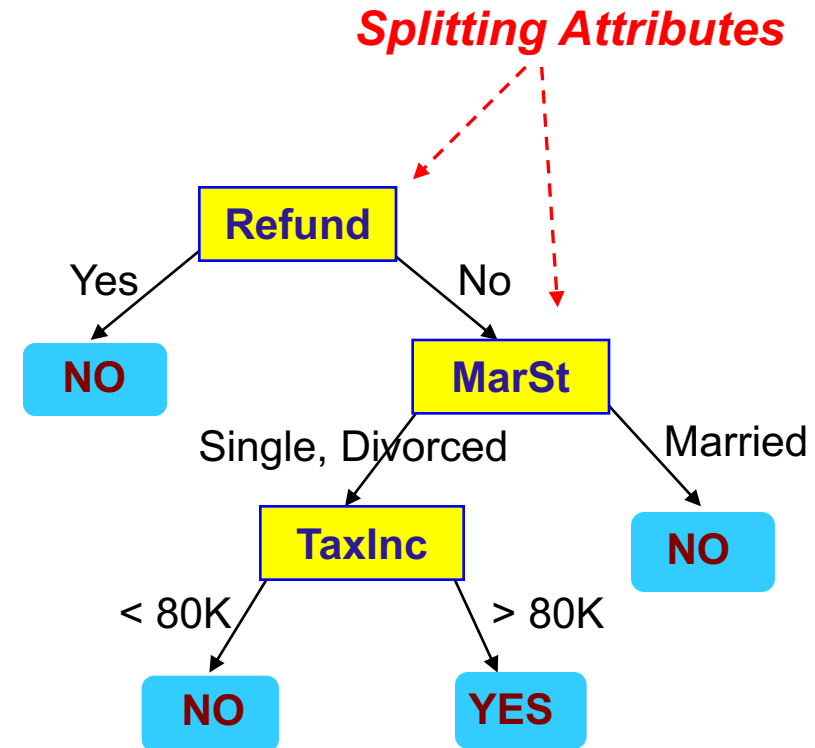
Number of Fisher Scoring iterations: 2



# EXAMPLE OF A DECISION TREE

<i>Tid</i>	<i>Refund</i>	<i>Marital Status</i>	<i>Taxable Income</i>	<i>Cheat</i>
1	Yes	Single	125K	No
2	No	Married	100K	No
3	No	Single	70K	No
4	Yes	Married	120K	No
5	No	Divorced	95K	Yes
6	No	Married	60K	No
7	Yes	Divorced	220K	No
8	No	Single	85K	Yes
9	No	Married	75K	No
10	No	Single	90K	Yes

*categorical*  
*categorical*  
*continuous*  
*class*



**Model: Decision Tree**

**Training Data**

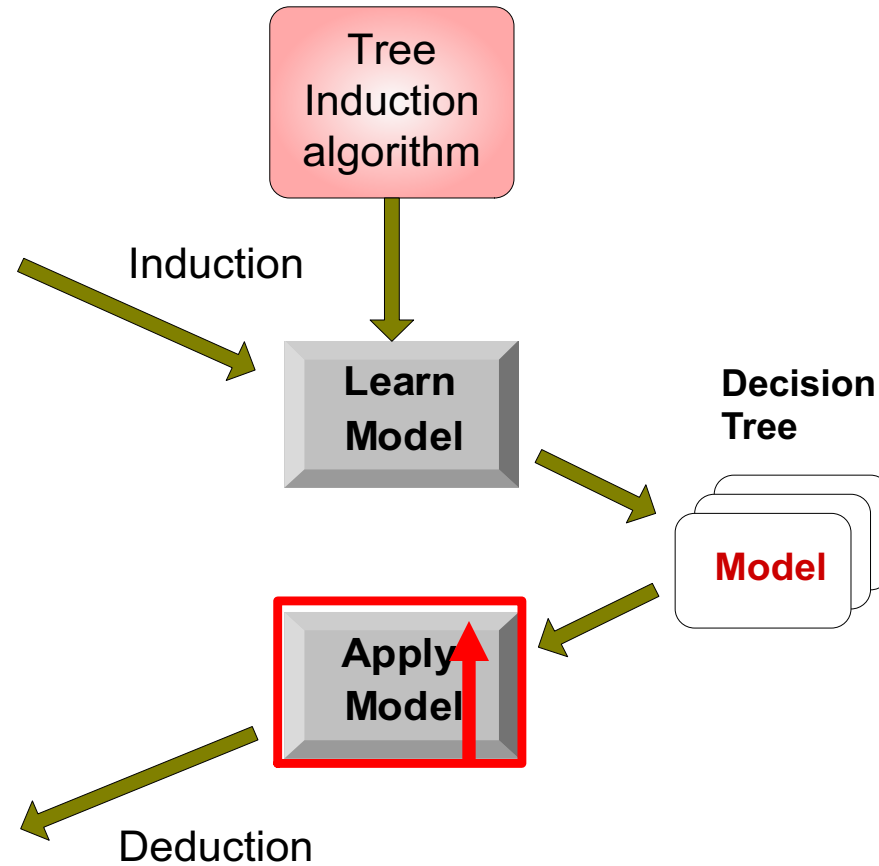
# DECISION TREE CLASSIFICATION TASK

Tid	Attrib1	Attrib2	Attrib3	Class
1	Yes	Large	125K	No
2	No	Medium	100K	No
3	No	Small	70K	No
4	Yes	Medium	120K	No
5	No	Large	95K	Yes
6	No	Medium	60K	No
7	Yes	Large	220K	No
8	No	Small	85K	Yes
9	No	Medium	75K	No
10	No	Small	90K	Yes

Training Set

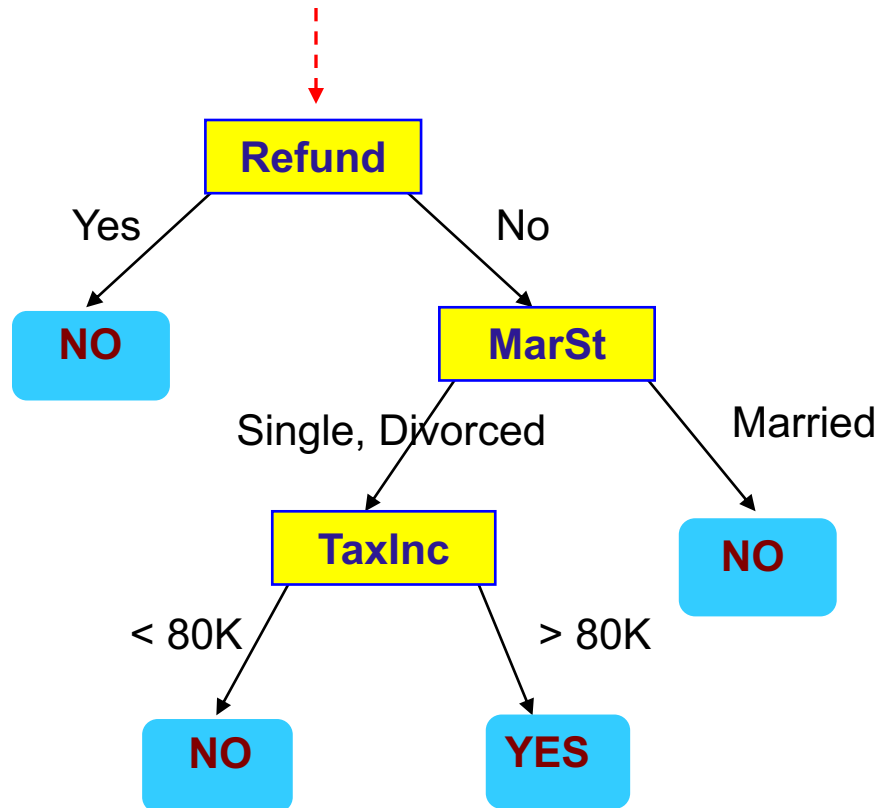
Tid	Attrib1	Attrib2	Attrib3	Class
11	No	Small	55K	?
12	Yes	Medium	80K	?
13	Yes	Large	110K	?
14	No	Small	95K	?
15	No	Large	67K	?

Test Set



# APPLY MODEL TO TEST DATA

Start from the root of tree.



## Test Data

Refund	Marital Status	Taxable Income	Cheat
No	Married	80K	?

# DECISION TREES

- Used for classifying data by partitioning attribute space
- Tries to find decision boundaries for specified optimality criteria
- Leaf nodes contain class labels, representing classification decisions
- Keeps splitting nodes based on split criterion, such as
  - GINI index, information gain or entropy
- Pruning necessary to avoid overfitting

# DECISION TREES IN R

```
mydata<-data.frame(iris)
attach(mydata)

library(rpart)

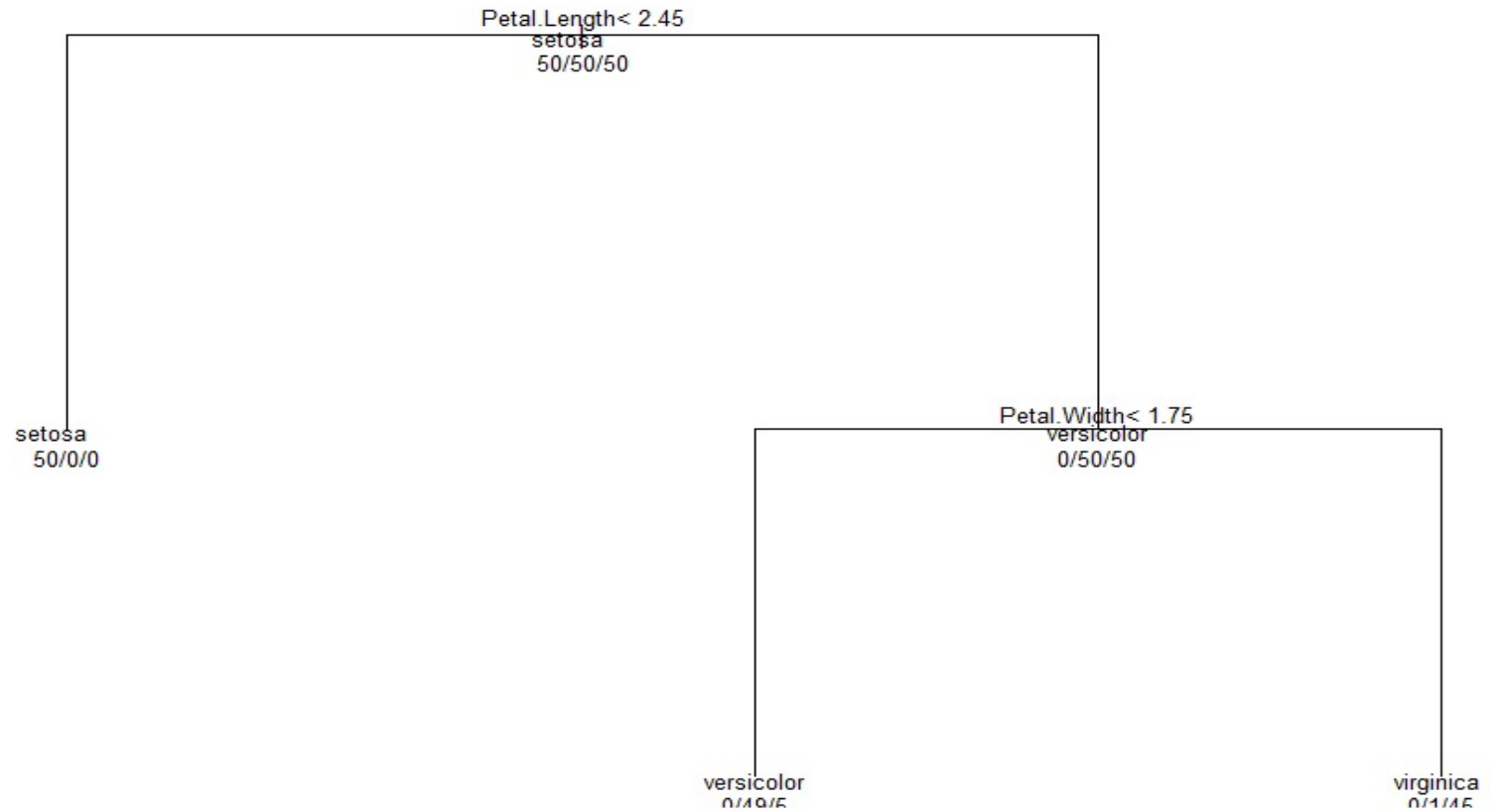
model<-rpart(Species ~ Sepal.Length +
  Sepal.Width + Petal.Length +
  Petal.Width,

  data=mydata,

  method="class")

plot(model)

text(model,use.n=TRUE,all=TRUE,cex=0.8)
```



# DECISION TREES IN R

```
library(tree)

modell<-tree(Species ~ Sepal.Length +
  Sepal.Width + Petal.Length +
  Petal.Width,

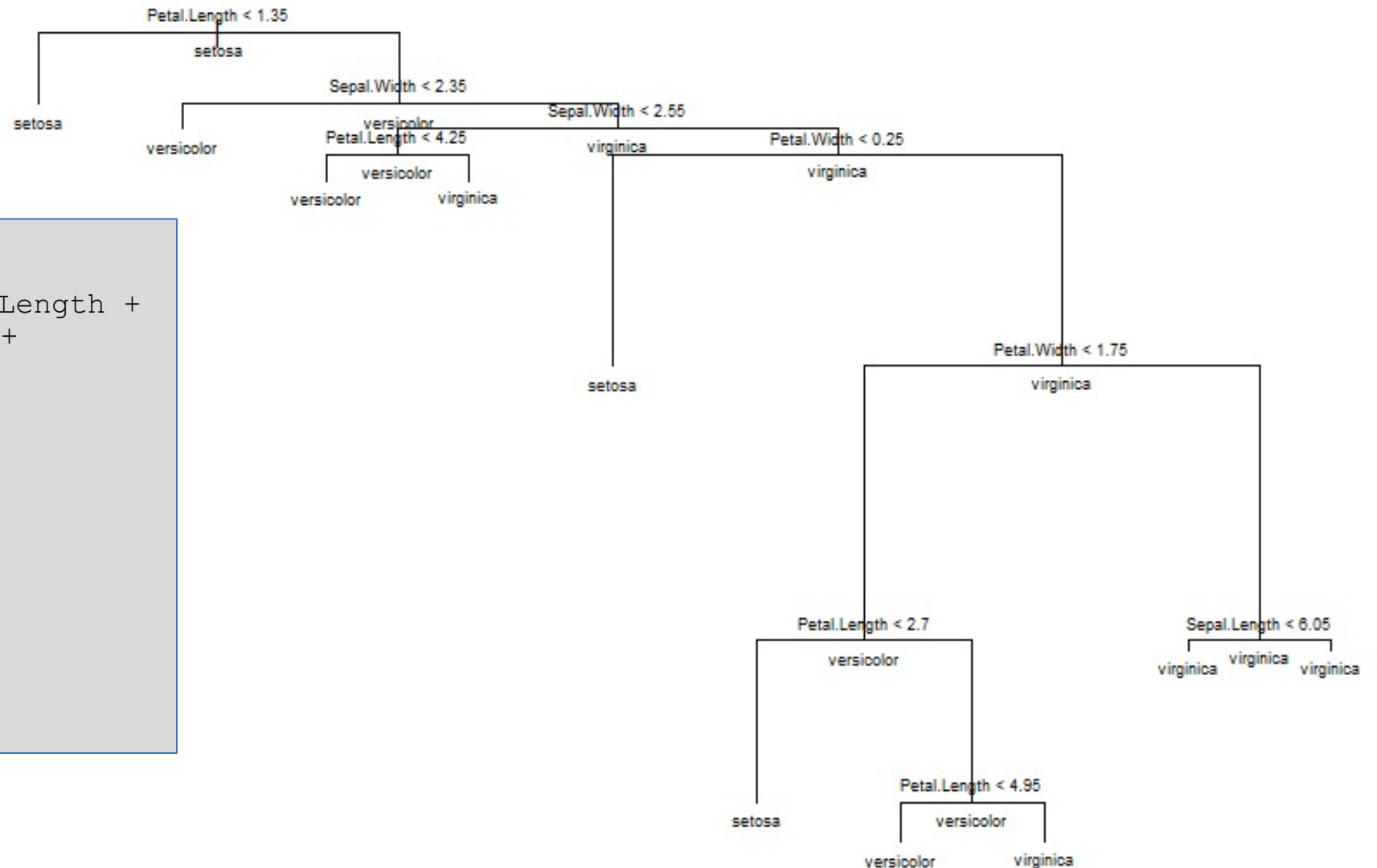
  data=mydata,

  method="class",

  split="gini")

plot(modell)

text(modell,all=TRUE,cex=0.6)
```



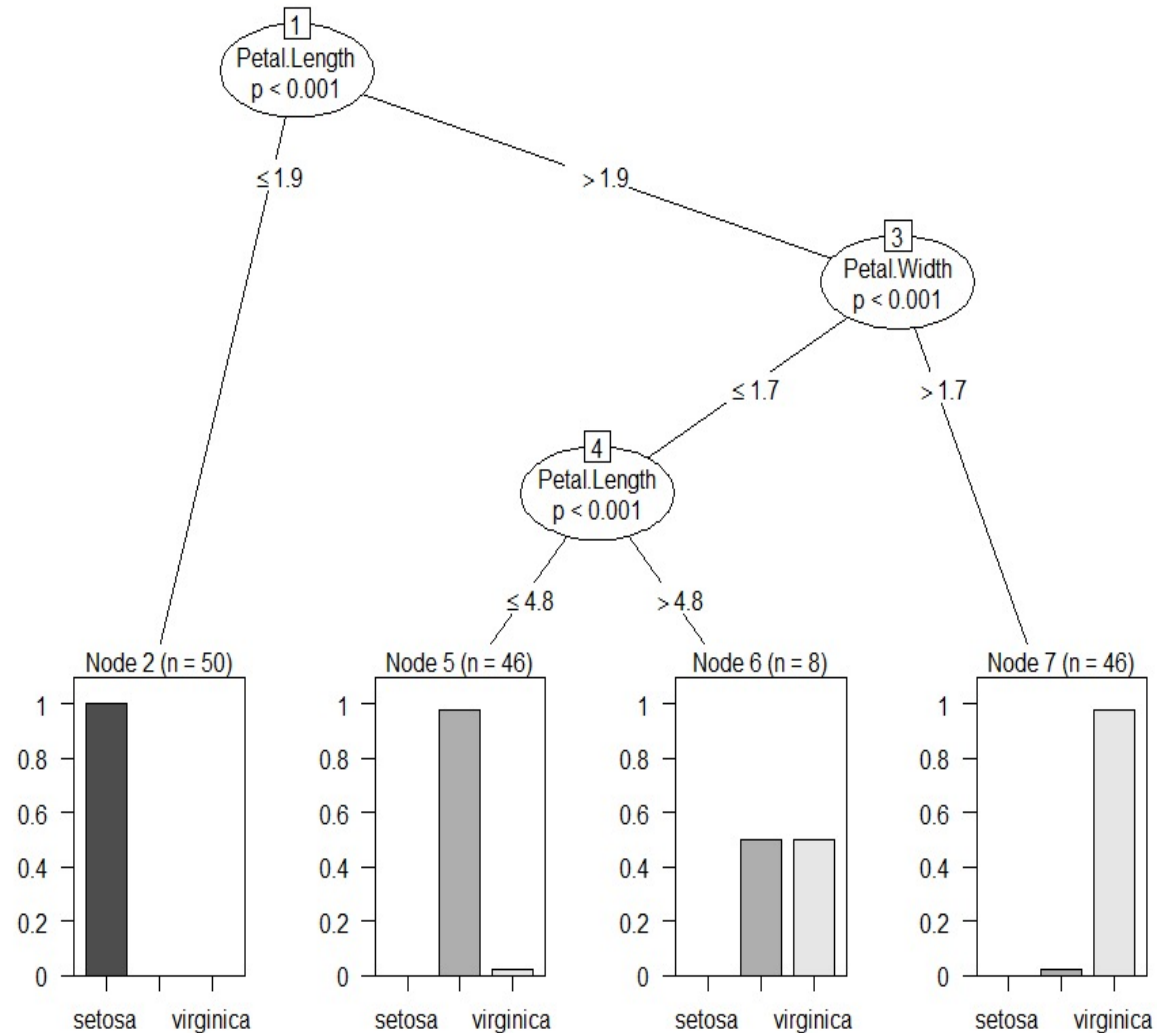
# DECISION TREES IN R

```
library(party)

model2<-ctree(Species ~
  Sepal.Length +
  Sepal.Width +
  Petal.Length +
  Petal.Width,

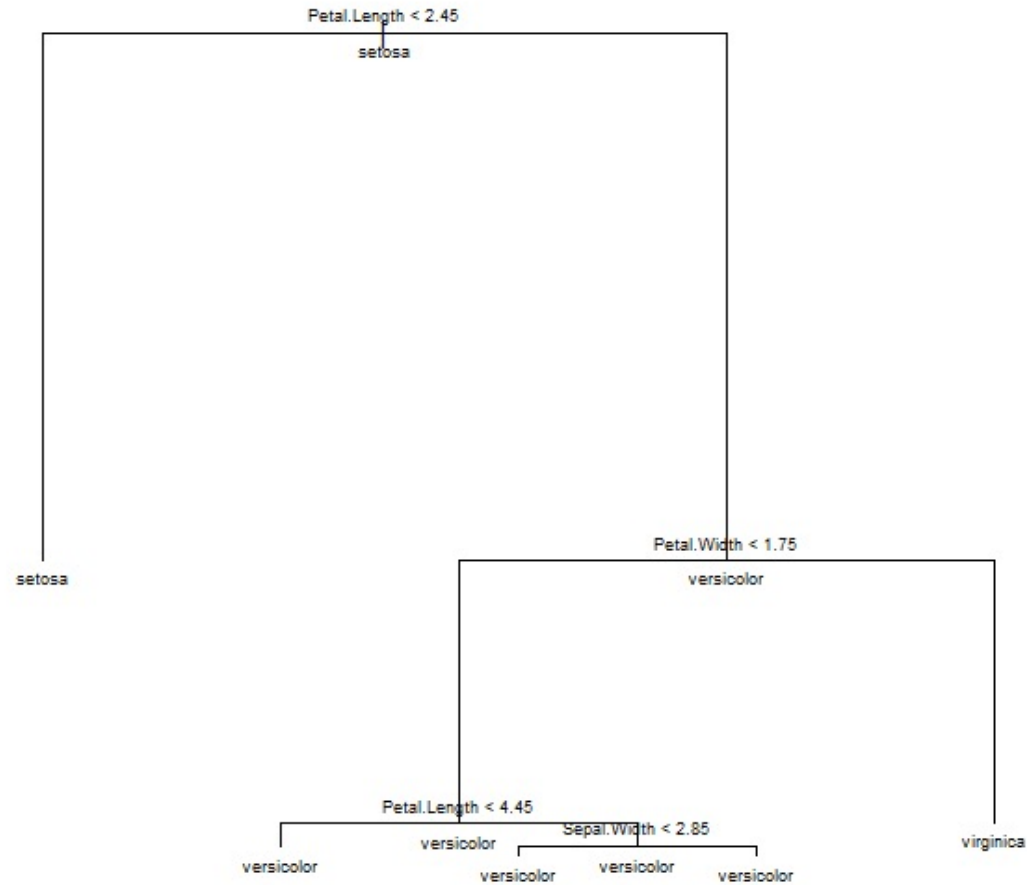
  data=mydata)

plot(model2)
```



# CONTROLLING NUMBER OF NODES

```
library(tree)
mydata<-data.frame(iris)
attach(mydata)
modell<-tree(Species ~ Sepal.Length +
Sepal.Width + Petal.Length +
Petal.Width,
  data=mydata,
  method="class",
  control = tree.control(nobs =
150, mincut = 10))
plot(modell)
text(modell,all=TRUE,cex=0.6)
predict(modell,iris)
```



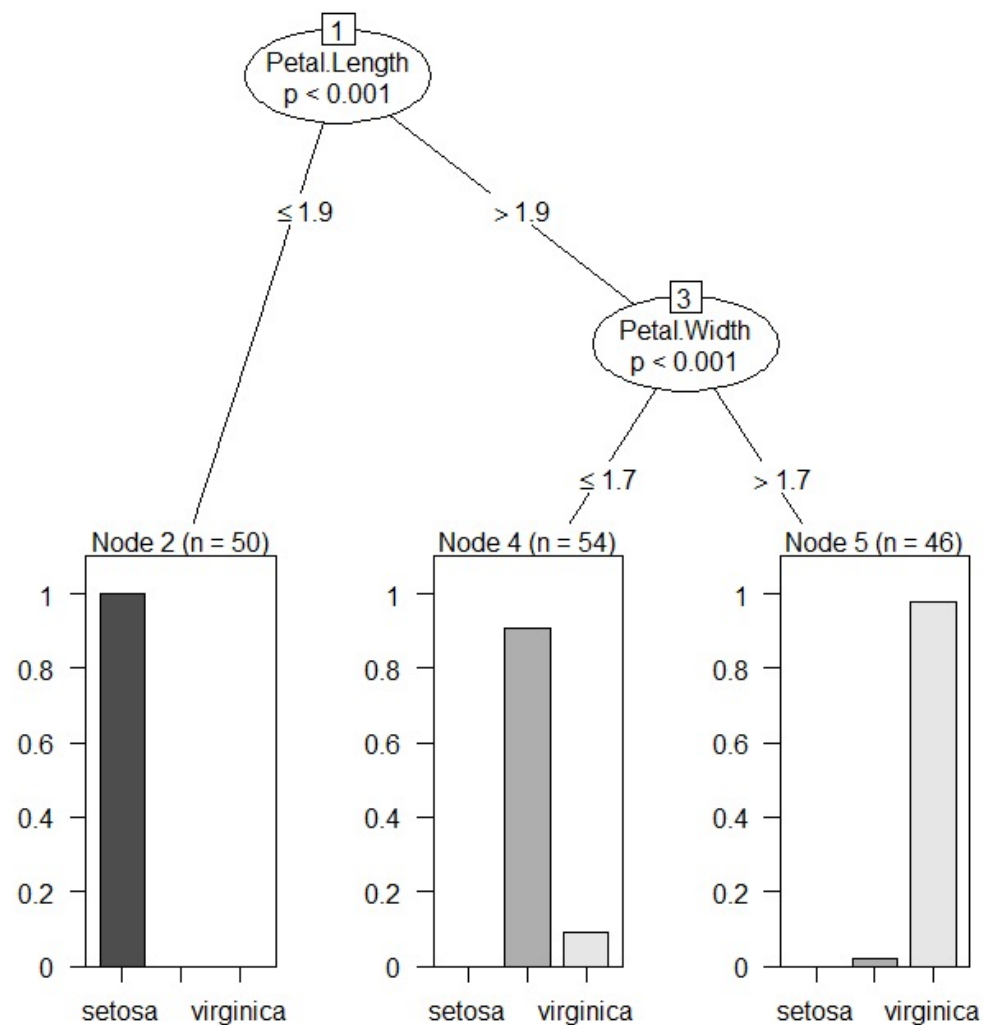
Note how the number of nodes is reduced by increasing the minimum number of observations in a child node!



# CONTROLLING NUMBER OF NODES

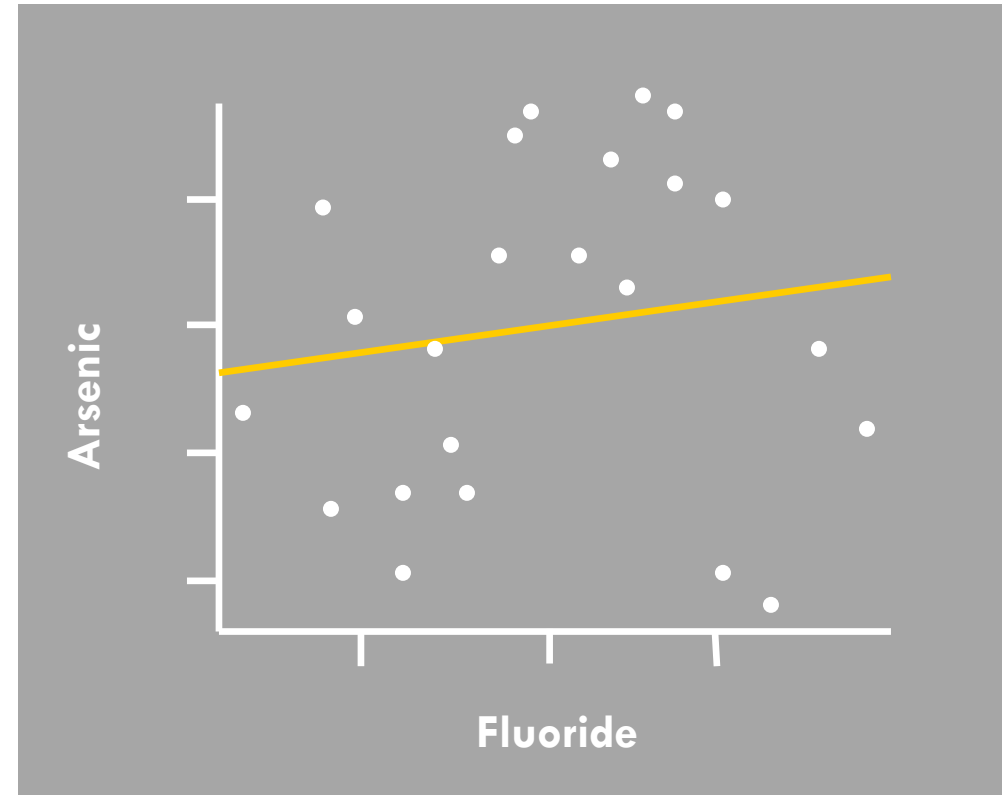
```
model2<-ctree(Species ~  
Sepal.Length + Sepal.Width +  
Petal.Length + Petal.Width,  
data = mydata, controls =  
ctree_control(maxdepth=2))  
  
plot(model2)
```

Note that setting the maximum depth to 2  
has reduced the number of nodes!



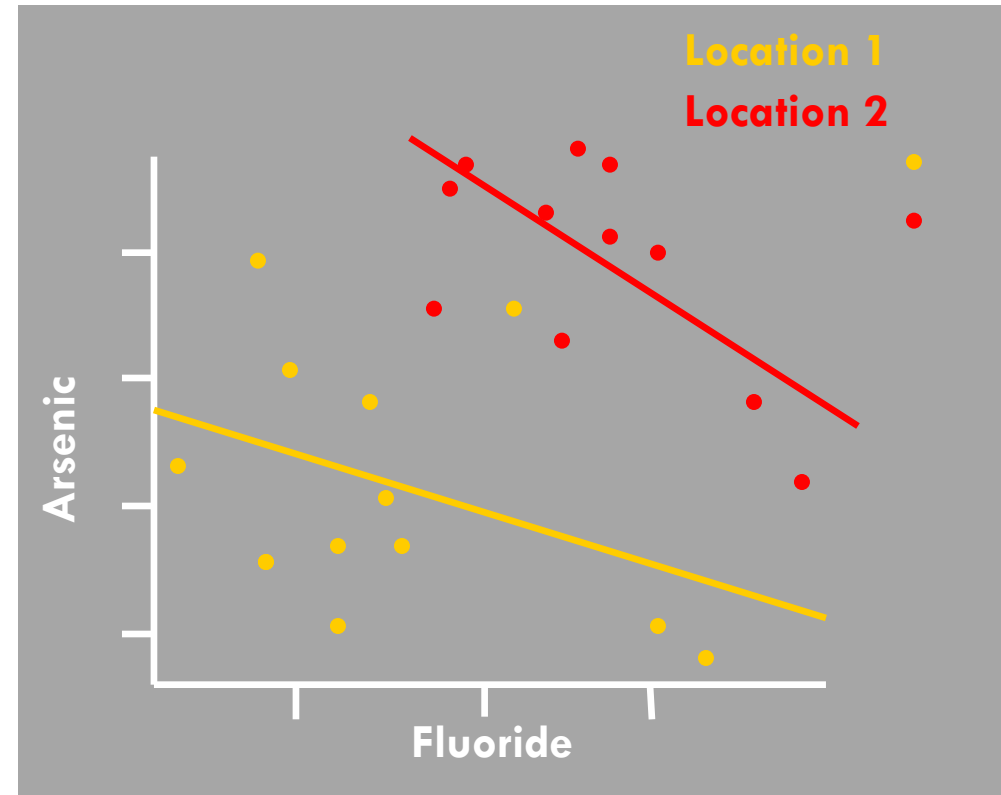
# SPATIAL VARIATION IN MODEL FIT

- When fitting a 'global' model (i.e., a single model) for a process observed over different spatial locations, we must assume that the relationship(s) are constant over space
  - this is often an incorrect assumption
- Here we have a line of best fit through two variables of water quality parameters, fluoride and arsenic



# SPATIAL VARIATION IN MODEL FIT

- Split up global data into regions and fit separate models for each region
- The challenge is how to define homogeneous regions
  - neighbourhoods
  - ecological zones
  - spatially-constrained cluster analysis
- The extreme is to estimate a new model at each location with a subset of neighbouring observations as the dataset
  - this is called 'geographically-weighted regression'



# KEY CONCEPTS TO CONSIDER WHEN WORKING WITH ENVIRONMENTAL MODELS

- **Data input:** quality, sources of bias, errors, consistency, training vs. testing
- **Model characteristics:** number of parameters, complexity, assumptions, representations
- **Model evaluation:** model fit, overfitting / generalizability
- **Model use:** how will models results be used, how can they be misused, etc.



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