

# Model Selection

- Philosophy of science and multiple alternative models
- Trade-offs
- Likelihood-based metrics
  - Likelihood Ratio Test
  - AIC
- Bayesian metrics
  - DIC
  - Predictive Loss

# “The” Scientific Method?

- Popper
  - Falsification of hypotheses
    - Hypotheses can not be proved, only disproved
- Stats: “Null hypothesis testing” (Fisher)
  - Single hypothesis is disproved by confrontation with the data
  - Likelihood the data would have been observed if the null hypothesis was true
  - If this probability (p-value) is small enough we reject the null

# Alternative Philosophies of Science

- Kuhn – Scientific Paradigms
  - Dominant paradigm used until there is so much contradictory information that it is “overthrown”
  - Requires an alternate paradigm that is “better”
- Polanyi – Republic of science
  - Multiple views of the world by different scientists
  - Confrontation between views and data judged by plausibility, value, and interest
- Lakatos – Scientific research program
  - Confrontation of multiple hypothesis with data as arbitrator

# Null models

- All these alternatives acknowledge
  - There may be multiple alternative models
  - Simple null models often scientifically trivial, uninteresting
  - Doesn't make sense to reject a model if there is not an alternative
- Likelihood and Bayesian stats both well suited to “judge” the contest between multiple competing hypotheses and data

# Models vs Hypotheses

- Models usually more specific than hypothesis
- Hypoth: Birds forage more efficiently in flocks
- Models: Consumption vs Size

- Consumption proportional

$$C = aS$$

- Consumption saturates

$$C = \frac{aS}{1 + bS}$$

- Increases then decreases

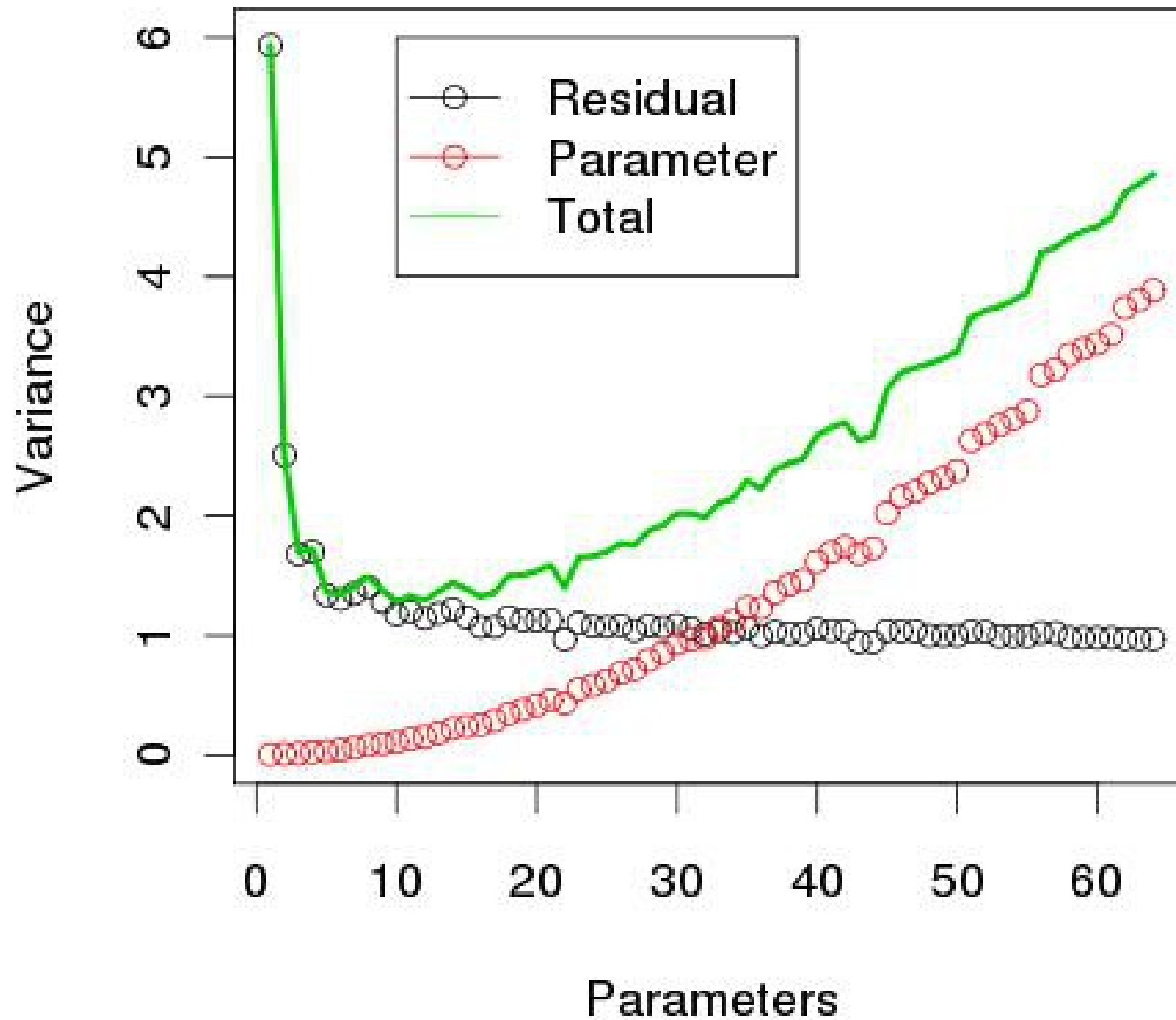
$$C = aS e^{-bS}$$

- “All models are wrong but some are useful”  
-- George Box

# Model selection

- Focus on choosing between multiple competing models rather than refuting a single null model
- How do we judge models?
  - Complexity
    - Number of parameters
  - Uncertainty
    - Model residuals
    - Parameter error (identifiability)
  - Data as ultimate arbiter
- “Make everything as simple as possible, but not simpler.” - A. Einstein

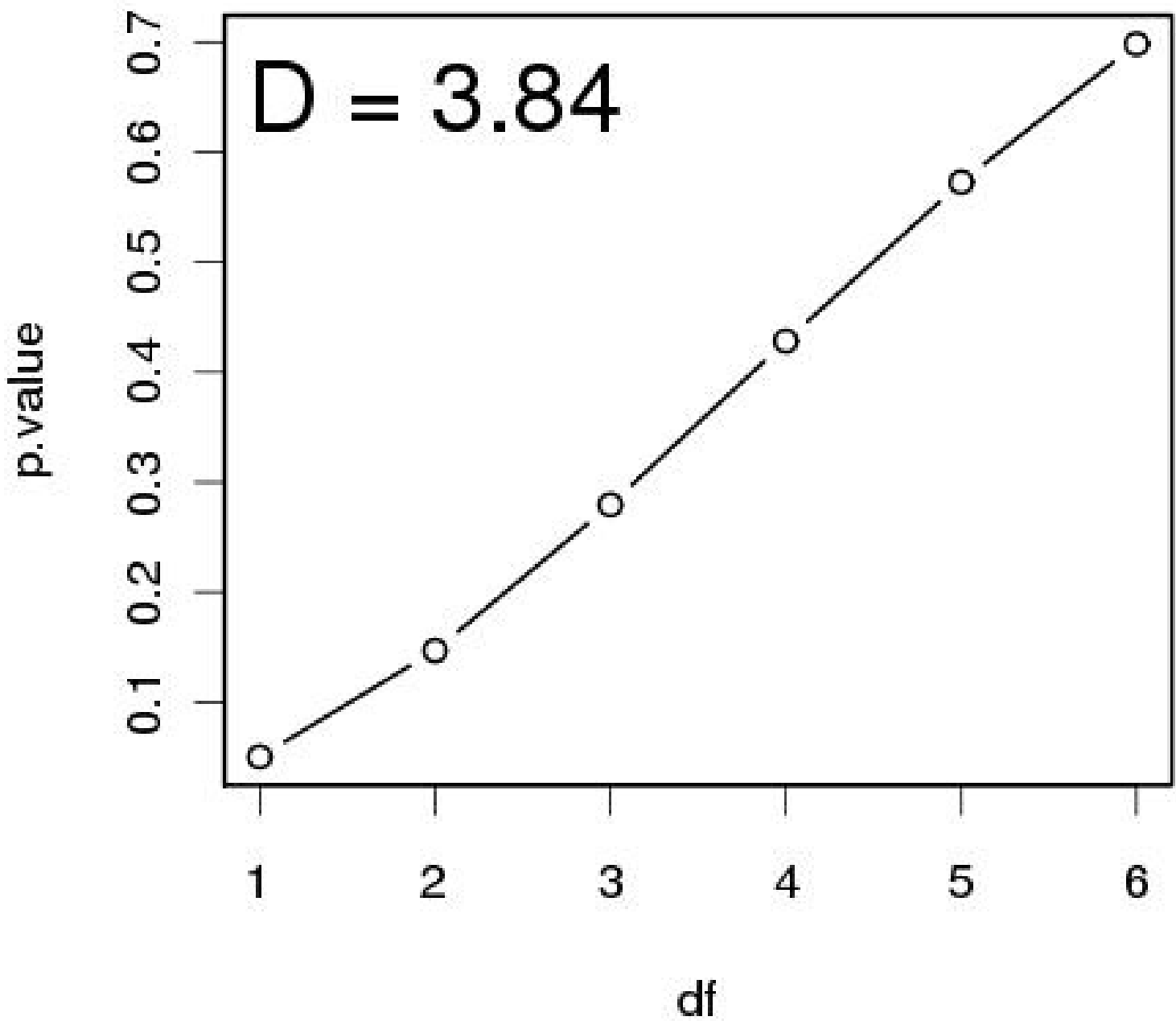
# UNCERTAINTY



# Likelihood Ratio Test

- $LR = L(x|\theta_0) / L(x|\theta_1)$
- $D = -2\ln L(x|\theta_0) - -2\ln L(x|\theta_1)$
- The test statistic D is known to be distributed with a  $\chi^2$  distribution
- Degrees of freedom = Difference in # of param.
  - Overall, L increases (-lnL declines) with # of param.
  - Penalizes model with more parameters
- $p\text{-val} = 1 - pchisq(D, df)$





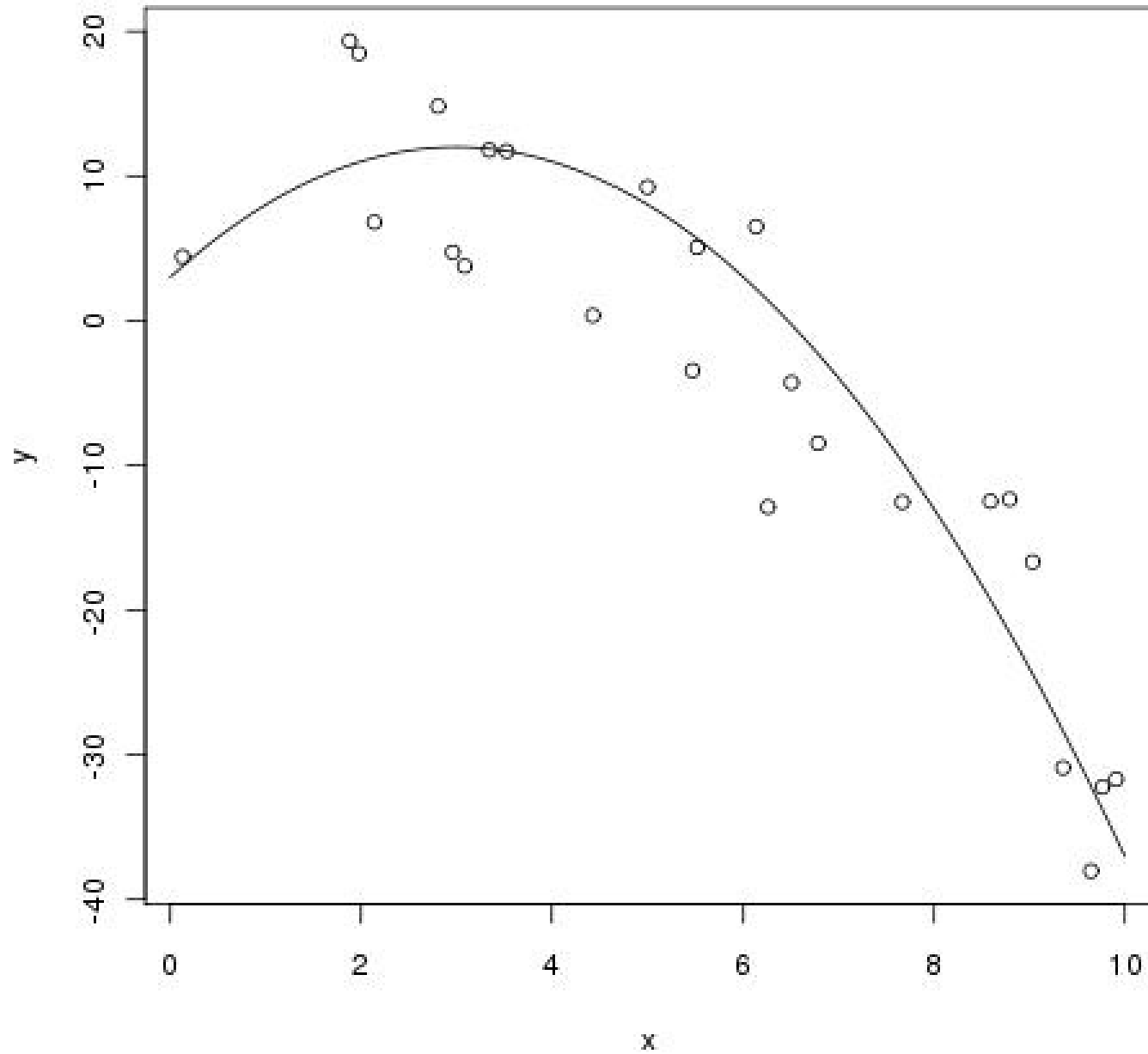
# LRT pro/con

- Only applies to nested models
- Asymptotically, slightly biased toward more complex models
- Provides a p-value
- Additional reminders:
  - **ALL** model selection criteria require application to the same data with same sample size
  - e.g. If adding covariate Z requires rows to be dropped because of missing values, have to drop from the model w/o Z as well

# Nested Models

- The more complex model collapses to the simpler model when one or more of the parameters is FIXED
- Examples:
  - Weibull vs Exponential (Lab 3)  
(fix  $c=1$ )
  - Pine cone: combined vs AMB/ELEV (Lab 4)
  - Regression: Inclusion of additional covariates  
(fix slope = 0)

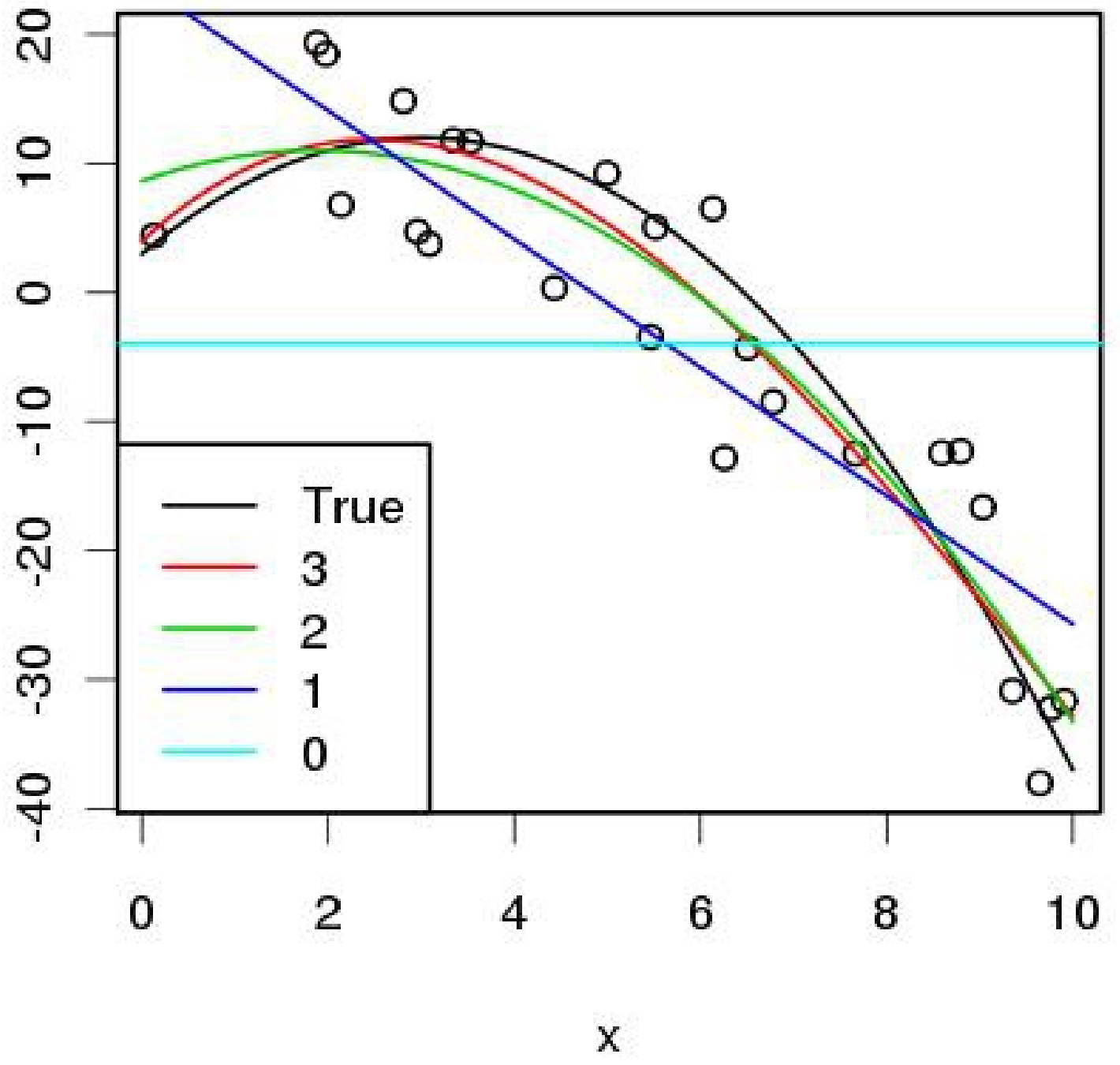
# Example: Polynomial



# Example: Polynomial

- Candidate models:
  - $Y = b_0$
  - $Y = b_0 + b_1 \cdot x$
  - $Y = b_0 + b_1 \cdot x + b_2 \cdot x^2$
  - $Y = b_0 + b_1 \cdot x + b_2 \cdot x^2 + b_3 \cdot x^3$
- Comparisons
  - 0 vs 1
  - 1 vs 2
  - 2 vs 3

- 0 vs 1  
 $p=7.6e-10$
- 1 vs 2  
 $p=0.00019$
- 2 vs 3  
 $p=0.9238$

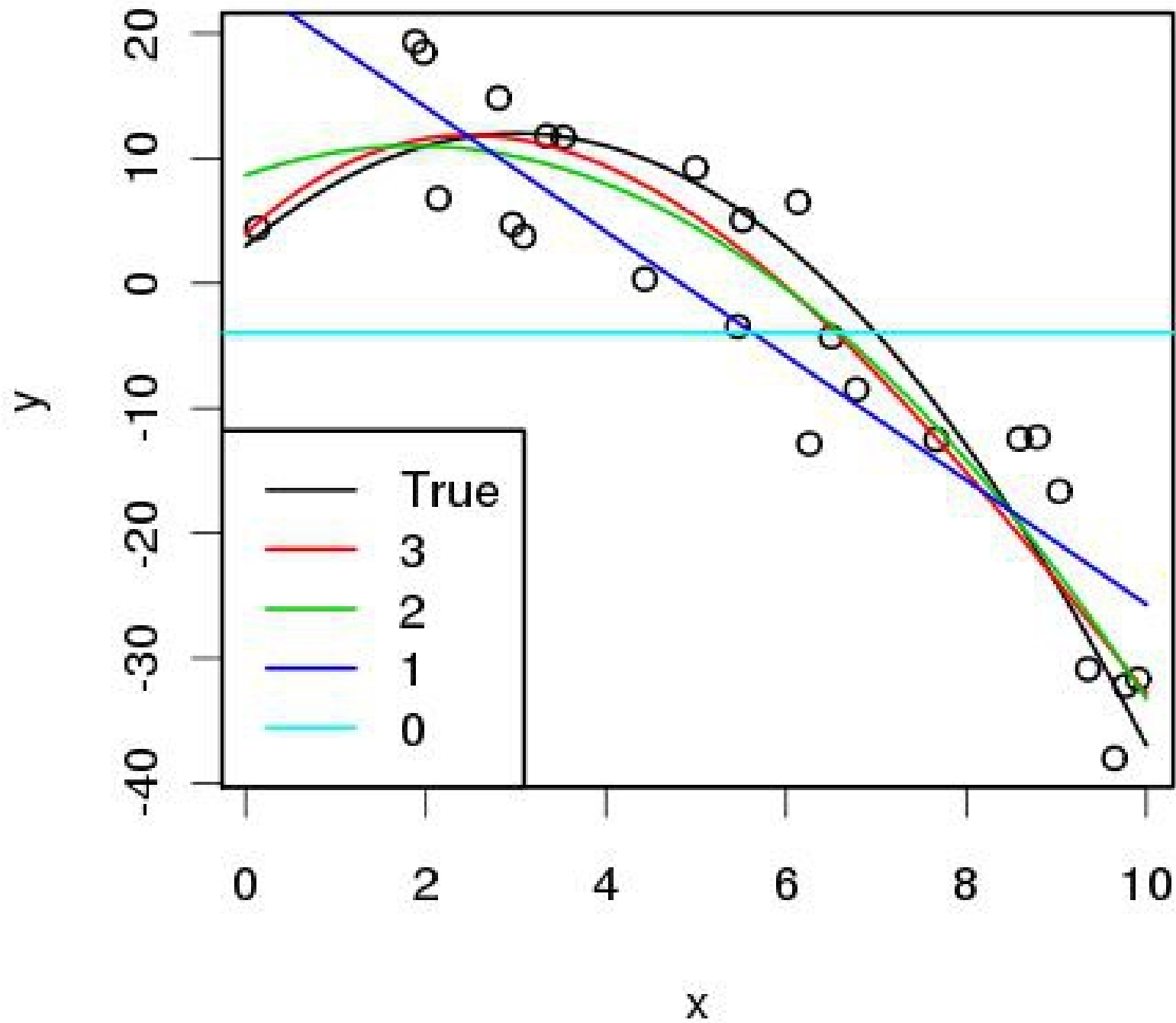


# Akaike Information Criterion

$$AIC = -2 \ln L + 2p$$

- $p$  = number of parameters in the model
- Based on information theory
- Lowest value “wins”
- Often expressed relative to best model,  $\Delta AIC$
- No p-value
- “Rules of thumb”
  - 0-2 = similar      2-5 = weak support      >5 = strong

- $\Delta AIC$
- 0  
47.77
- 1  
11.91
- 2  
0.00
- 3  
1.99





# P-value

- Probability of obtaining a test statistic at least as extreme as the one that was actually observed, assuming that the null hypothesis is true.
- **Not** the probability that the null hypothesis is true
  - P-value can be close to zero when the posterior probability of the null is close to 1
- **Not** the probability of falsely rejecting the null hypothesis

# PROBABLE CAUSE

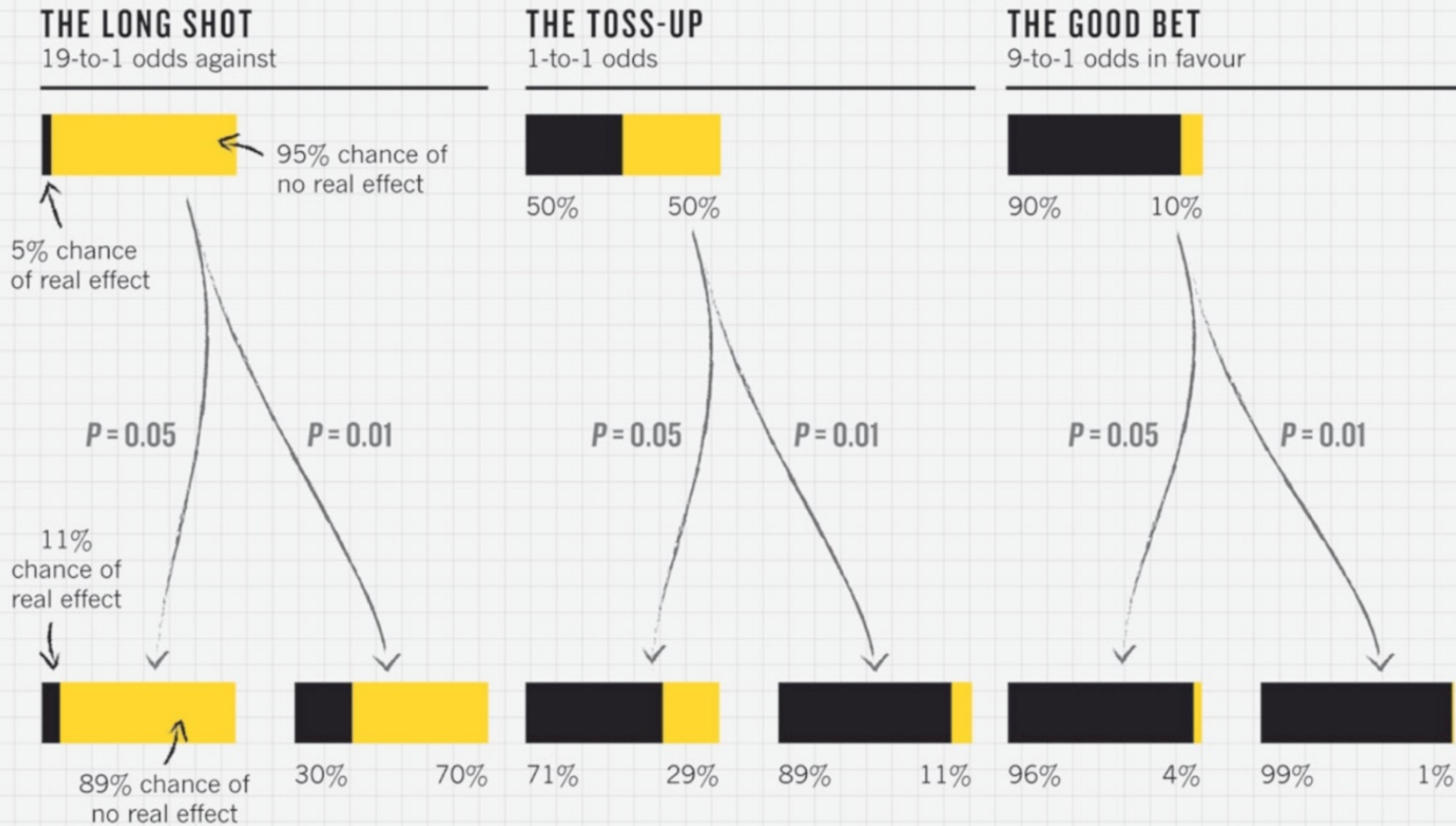
A  $P$  value measures whether an observed result can be attributed to chance. But it cannot answer a researcher's real question: what are the odds that a hypothesis is correct? Those odds depend on how strong the result was and, most importantly, on how plausible the hypothesis is in the first place.

■ Chance of real effect  
 ■ Chance of no real effect

**Before the experiment**  
 The plausibility of the hypothesis — the odds of it being true — can be estimated from previous experiments, conjectured mechanisms and other expert knowledge. Three examples are shown here.

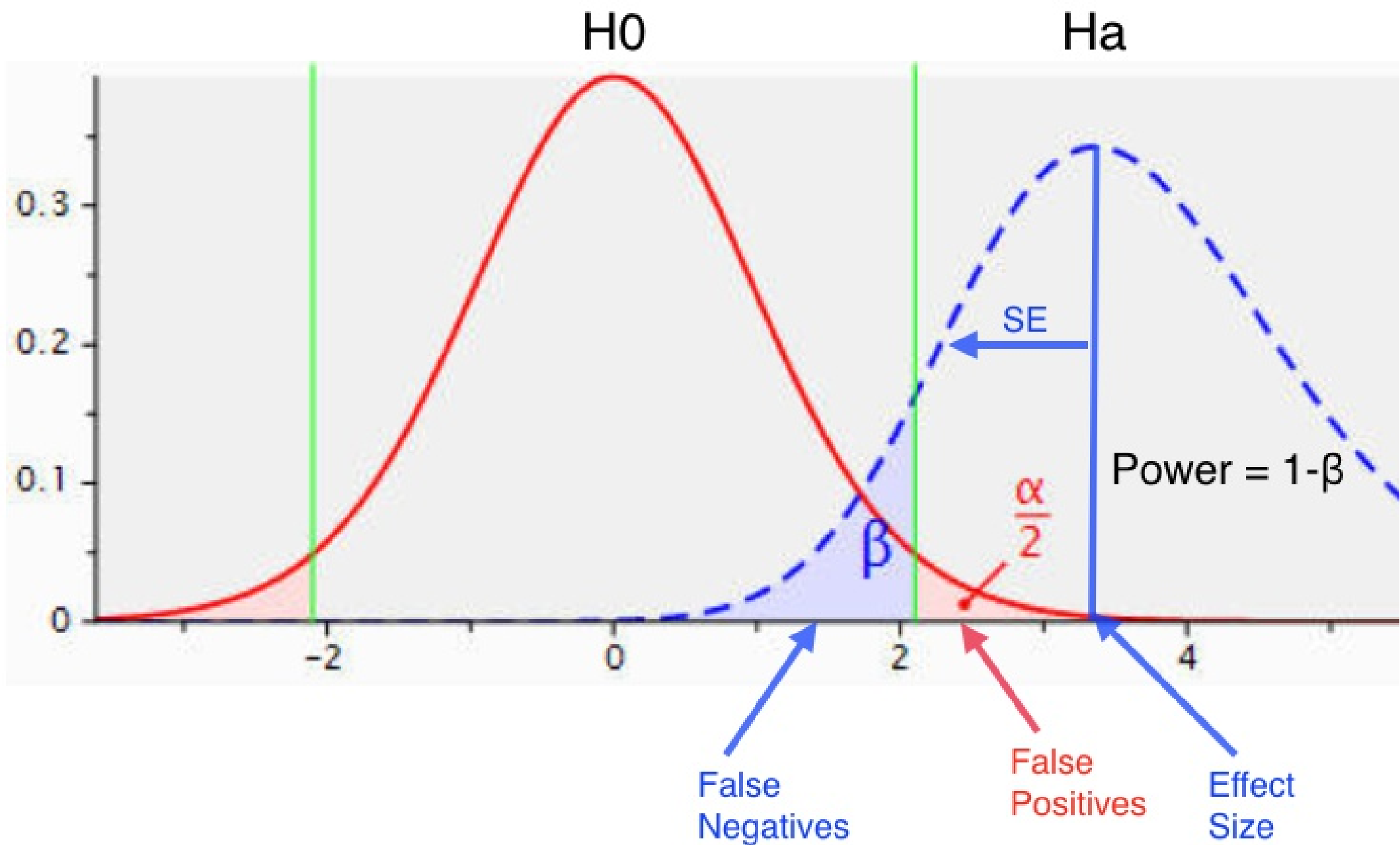
**The measured  $P$  value**  
 A value of 0.05 is conventionally deemed 'statistically significant'; a value of 0.01 is considered 'very significant'.

**After the experiment**  
 A small  $P$  value can make a hypothesis more plausible, but the difference may not be dramatic.



# Example: Southern Brown Frog

- Researcher surveys a pond for the frog
- From prior experience 80% detection | present
- No frogs observed
- If null hypothesis is frogs are absent
  - $P = 1.0$  -- Fail to reject
  - Further surveys that fail to find the frog,  $p=1.0$
- If null hypothesis is frogs are present
  - $P = 0.2$  – Fail to reject



Power = f(effect size, SE)

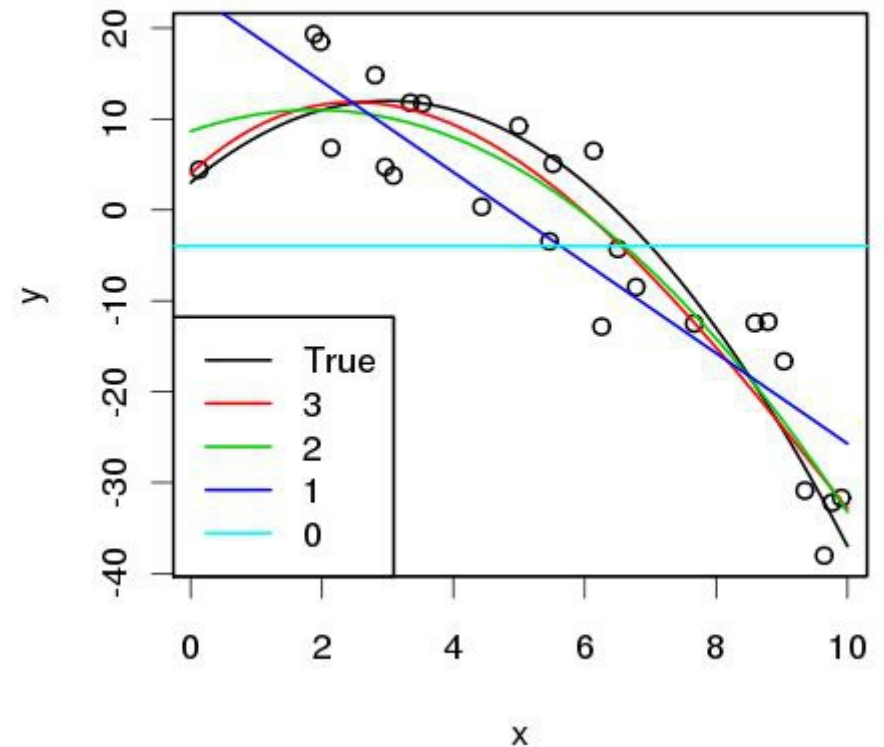
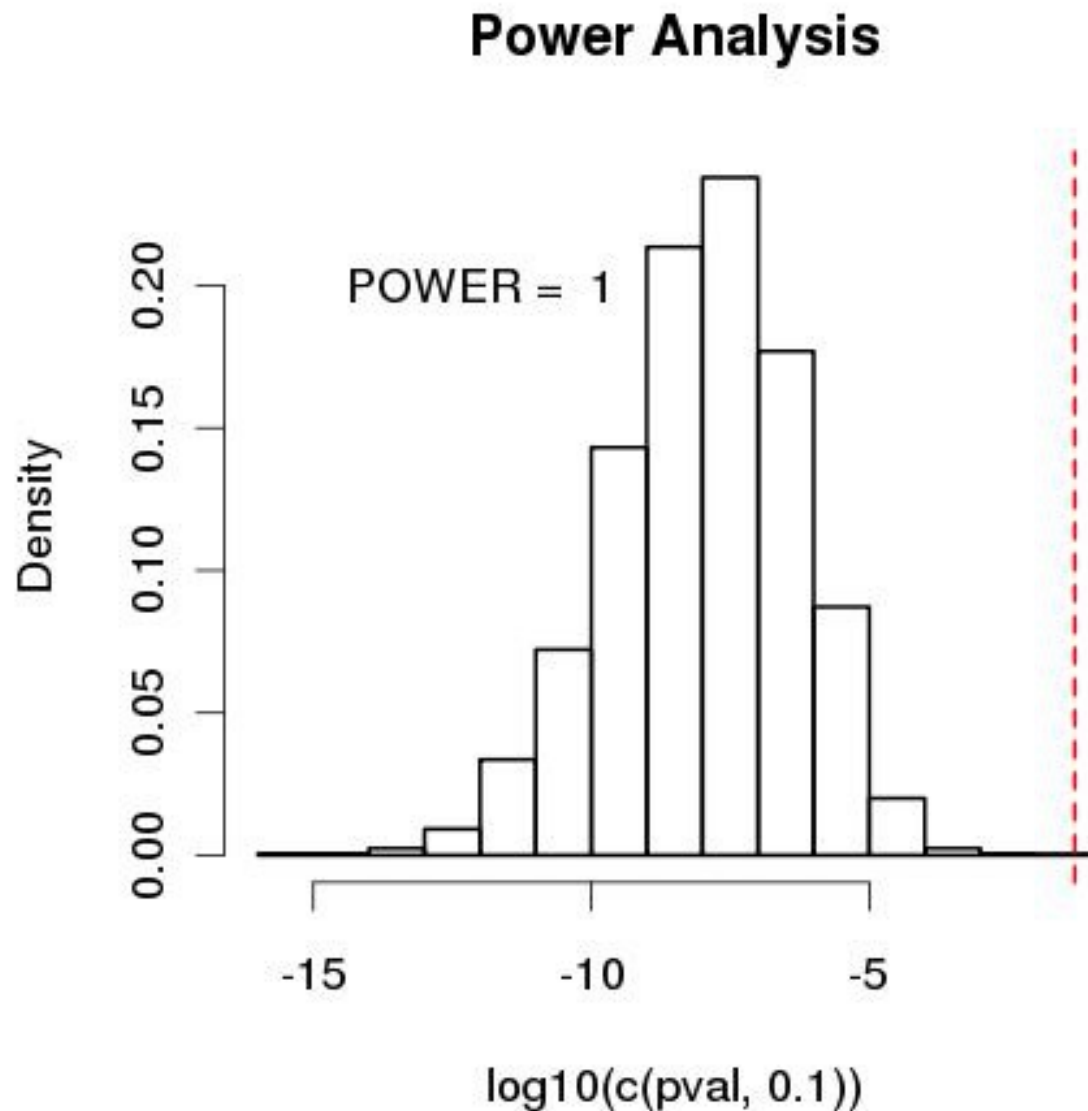
# Power

- Probability of correctly rejecting the null hypothesis
- Requires that some explicit alternative hypothesis is stated
  - Parameter values
  - Variance
  - Sample size
- Often calculated as a function of sample size
- For complex models, calculate through simulation

# Generic Example

```
LnL.A = function(theta){
  -sum(dnorm(y,f(x,theta),sd)))
}
lnL.0 = function(mu){
  -sum(dnorm(y,mu,sd))
}
for(i in 1:nsim){
  Ey = f(x,theta)      ## process model
  y = rnorm(N,Ey,sd)   ## data model
  outA = optim(ic,lnL.A) ##fit of alternative
  out0 = optim(ic,lnL.0) ##fit of null
  pval[i] = 1-pchisq(2*(outA$value-out0$value),df)
}
power = sum(pval < 0.05)/nsim
```

# Example: Quadratic vs Linear LRT



- Results **specific** to parameter values and sample size chosen

# Identifiability

- Data may not provide information on all parameters in a model
- Often requires restructuring model
- Not fixed by collecting more data
- Parameters often “trade-off when fitting”
- Simple examples
  - $N(\mu, \sigma^2 + \tau^2)$
  - $N(a/b, \sigma^2)$
- Occur in both Likelihood and Bayes