Regression and Modeling in R

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1. Regression: Biomass data

Use log-transformed data Possible terms

- a) simple model: log(AGB) vs. log(dbh)
- b) second order term: log(dbh) squared
- c) rainfall and elevation

Graph

- a) log(AGB) vs. log(dbh)
- b) add curve of best fit
- c) overlay curves for high rainfall and low rainfall

Character variable (factors)

- a) use ForestType in the model
 - $\log(volume) \sim \log(dbh) + ForestType$
- b) compare to 3 independent models (3 forest types)
 - $\log(volume[dry]) \sim \log(dbh[dry])$
 - etc.

2. Regression: Function to calculate linear regression

The basics

- a) Accept *x* and *y* values
- b) Return regression parameters
- c) Graph *y* vs. *x*
- d) Adds regression line

More information

- a) Find predicted *y* for every *x*
- b) Find residuals and their variance
- c) Graph residuals

Advanced (extra credit)

- a) Include option of logarithmic transformation
- b) Allow multiple *x* predictors
- c) Graph y as a function of a chosen x

3. Simulation: Create a simulated correlation and test how well Im fits the parameters

The basics

- a) Define *x* from normal distribution
- b) Define slope and intercept parameters
- c) Define error with *rnorm* and *sd*
- d) Calculate y
- e) Use *lm* to estimate slope and intercept

More information

- a) Evaluate impact of increasing error
- b) Evaluate impact of error in measuring x
- c) Evaluate impact of highly non-Gaussian *x*

Advanced (extra credit)

- a) Test multiple regression, with x_1 and x_2 predictors
- b) Evaluate impact of correlation between x_1 and x_2

4. Non-linear model

- Use optim to fit a model $y \sim x + x^2$, or $y \sim x + \log x$
- Fit a model to tree height data (dataset treeht)
- Consider this function of height as a function of dbh:
- $h = H_{max}(1 e^{-ax^b})$ where H_{max} , a, and b are parameters, h is height and x is dbh. Use data from a single species to estimate the 3 parameters using maximum likelihood and a Gaussian error.

5. Error functions

- dlnorm
 - for abundances, whether integer or not
 - good match for tree growth rates
 - but cannot handle zeroes
- dpois including zeroes (but does not handle most ecological data well)
 - for integer abundances
 - handles zeroes
 - however, close to Gaussian so not appropriate for much ecological data
- dnbinom (common in ecology)
 - for integer abundances that are highly skewed
 - very common in ecology
 - R: prob=dnbinom(count,size=k,mu=mu)
 - size is 'clumping parameter'; mu is mean

6. Two-parameter survival model

- Fit tree survival to growth and dbh
- $logit(status_2) \sim gr_1 + dbh_1$

7. Program a Gibbs sampler for tree height model

- Write the sampler for the 3-parameter height model (formula in assignment 3)
- Extend the program to allow a linear *height* ~ *dbh* model
- I provide a head start (*modelGibbs.r*)