

LINEAR MODELS	GENERALIZED LINEAR MODELS
Model statement:	Model statement:
$y_i = \beta_0 + \beta_1 \ x_{i1} + \beta_2 \ x_{i2} + \ldots + \beta_m \ x_{im} + \epsilon_i$ $\textbf{Continuous} \ response \ variable \ (normal \ distribution)$ $Predictor \ variables \ can \ be \ continuous \ and/or \ categorical$ $Can \ have \ transformations \ of \ x's \ and/or \ y$	$y_i = \mathbf{g^{-1}}(\beta_1 \ x_{i1} + + \beta_m \ x_{im}) + \epsilon_i$ Response variable is typically <b>binary</b> (from binomial distribution) or <b>count</b> (from Poisson distribution)  Response probability distributions can be any of the exponential family of distributions (e.g.: normal, inverse normal, gamma, binomial, negative binomial, Poisson,
to satisfy assumptions	multinomial)
Link function is identity link, with normal distribution (if you used a GLM to fit this)	Predictor variables can be continuous and/or categorical  Non-linear <b>link function</b> relates predictor variables and response variable (logit or probit link for binomial, log link for Poisson)
How model is fit	How model is fit
Least squares or maximum likelihood Least squares: find parameter estimates such that the sum of squared differences between errors is the smallest (minimize SSE)	Maximum likelihood: find parameter estimates such that the likelihood of getting these y-values is the largest (maximize likelihood)
Features	Features
Goes through the mean of all the x values and the y values Sum of errors is zero; $\Sigma e_i = 0$	<ol> <li>Error structure</li> <li>Link function</li> <li>Linear predictor</li> </ol>

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Assumptions	Assumptions
<ol> <li>Linearity between continuous predictor variables and response variable (residual plot)</li> <li>errors are independent and identically distributed (equal variance/ homoskedastic, no correlation between errors due to space/time/etc.) and follow the normal distribution (residual plot, heteroskedasticity tests, normality plot, normality tests, histogram of errors)</li> </ol>	<ol> <li>Variances of y's is a function of the mean of y, given x (residual plot can be used to check for overdispersion)</li> <li>Observations (errors) are independent</li> </ol>
<ul> <li>Sampling assumptions</li> <li>1. Predictor variables are measured without error</li> <li>2. Data represents a random sample from the population</li> </ul>	<ul> <li>Sampling assumptions</li> <li>1. Predictor variables are measured without error</li> <li>2. Data represents a random sample from the population</li> </ul>
Hypothesis tests	Hypothesis tests
F test comparing full model to null model  t tests to test contribution of each variable, given the other variables (equivalent to a partial F test* for that variable)  *partial F tests can be used to test multiple variables at the same time by comparing nested models	Likelihood ratio test comparing full model to null model  Likelihood ratio test comparing models with and without the variable of interest (can compare any set of nested models, therefore can test multiple variables at the same time)

### LINEAR MODELS

## Measures of goodness of fit

Maximize

$$1. \mathbf{R}^2 = SS \text{ reg} / SSY = 1 - SSE/SSY$$

2. 
$$\mathbf{R}^2$$
 adjusted =  $1 - (\underline{n-1})$  SSE  $\underline{\qquad}$  SSY

If y was transformed:

You will need to backtransform the predicted values into the original units. You can then calculate a Pseudo-R<sup>2</sup>:

$$Pseudo-R^2 = 1 - SSE/SSY$$

#### Minimize

- 1. **Root MSE** / Standard error of the estimate
- 2. PRESS (Prediction sum of squares) statistic subset used to estimate parameters, then tested at prediction for remaining data

## 3. Akaike's information criterion

(based on least squares)

 $AIC = n \ln(SSE) - n\ln(n) + 2p$ 

p = number of parameters

n = sample size

### GENERALIZED LINEAR MODELS

### Measures of goodness of fit

Maximize

Pseudo R<sup>2</sup> = 1 - 
$$\left\{ \frac{L(0)}{L(\hat{\beta})} \right\}^{\frac{2}{n}}$$

1.

L(0) = likelihood of intercept only model

### 2. Max-scaled $R^2$

$$\widetilde{R}^2 = \frac{R^2}{R^2 \max}$$

where 
$$R^2 \max = 1 - \{L(0)\}^{2/n}$$

### Minimize

# 1. Akaike's information criterion

$$AIC = -2\ln L + 2(k+s)$$

Where k =levels for y - 1, and s is the number of predictor variables (including all dummy variables).

AIC with correction for finite sample sizes

$$AICc = AIC + \frac{2k(k+1)}{n-k-1}$$

n = sample size

Penalty for extra parameters

AICc converges to AIC when n is large

### 2. Swartz's criterion (BIC)

$$SC = -2 \ln L + (k + s) \ln(\sum_{j=1}^{s} f_j)$$

 $f_i$  is the frequency for each proportion;

sum of frequencies = n when single observations of y = 0 or 1 are used.

Includes sample size in the form of frequencies

Also see **classification table** (results for different probability cut-offs) for logistic regression