## Influence & Transformations

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Readings: ALR 8-9, Gelman & Hill Ch 2-4

# Assumptions of Linear Regression



(inexv

$$Y_i = \beta_0 + \beta_1 X_{i1} + \beta_2 X_{i2} + \dots \beta_p X_{ip} + \epsilon_i$$

- $\blacktriangleright$  Model Linear in  $X_i$  but  $X_i$  could be a transformation of the original variables
- $ightharpoonup \epsilon_i \sim N(0, \sigma^2)$
- - correct mean function
  - constant variance

 $Y \mid X, \beta, \sigma$ ► independent Normal errors Wormal linearity transform of x

#### Animals

Read in Animal data from MASS. The data set contains measurements on body weight (kg) and brain weight (g) on 28 animals.

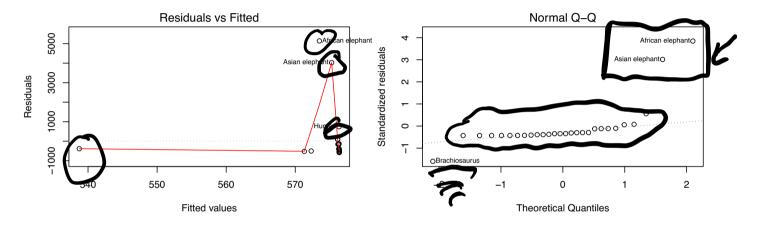
Let's try to predict brain weight from body weight.

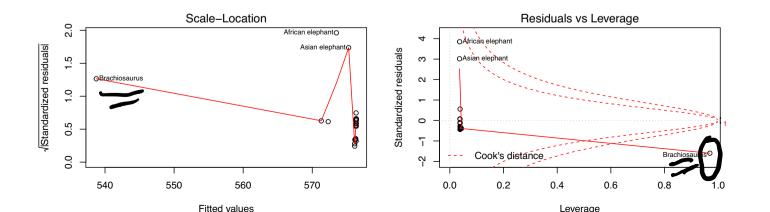
```
data(Animals, package="MASS")
brain.lm = lm(brain ~ body, data=Animals)
```

### Diagnostic Plots

## Warning in sqrt(crit \* p \* (1 - hh)/hh): NaNs produced

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#### **Outliers**

▶ Flag outliers after Bonferroni Correction  $p_i < \alpha/n$ 

```
pval = 2*(1 - pt(abs(rstudent(brain.lm)), brain.lm$df -1))
rownames(Animals)[pval < .05/nrow(Animals)]</pre>
```

```
## [1] "Asian elephant" "African elephant"
```

 Use functions from the CAR package (Companion to Applied Regression) for Bonferroni correction

```
car::outlierTest(brain.lm)
```

```
## African elephant 5.751645 5.4098e-06 0.0001514
## Asian elephant 3.667458 1.1576e-03 0.0324120
```

Measure of influence of case i on predictions

$$D_i = \frac{\|\hat{\mathbf{Y}} - \hat{\mathbf{Y}}_{(i)}\|^2}{\hat{\sigma}^2 \, p}$$

after removing the ith case

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Easier way to calculate

$$D_i = \frac{e_i^2}{\hat{\sigma}^2 p} \left[ \frac{h_{ii}}{(1 - h_{ii})^2} \right],$$

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Easier way to calculate

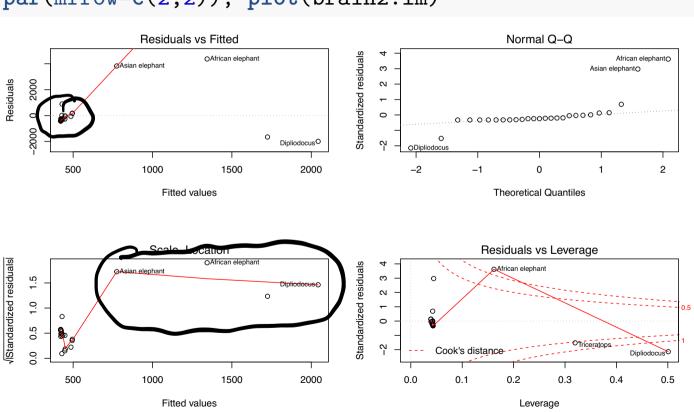
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ight],$$
  $D_i = rac{r_{ii}}{p} rac{h_{ii}}{1 - h_{ii}}$ 

Flag cases where  $D_i > 1$  or  $D_i > 4/n$ 

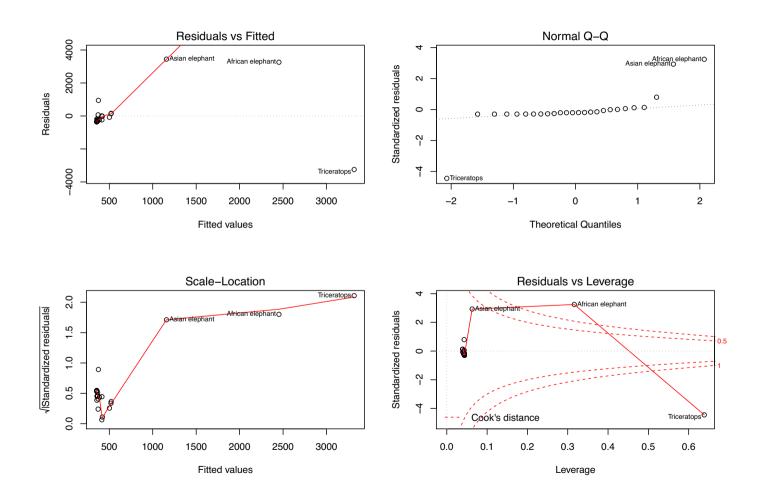
rownames(Animals)[cooks.distance(brain.lm) > 1]

## [1] "Brachiosaurus"

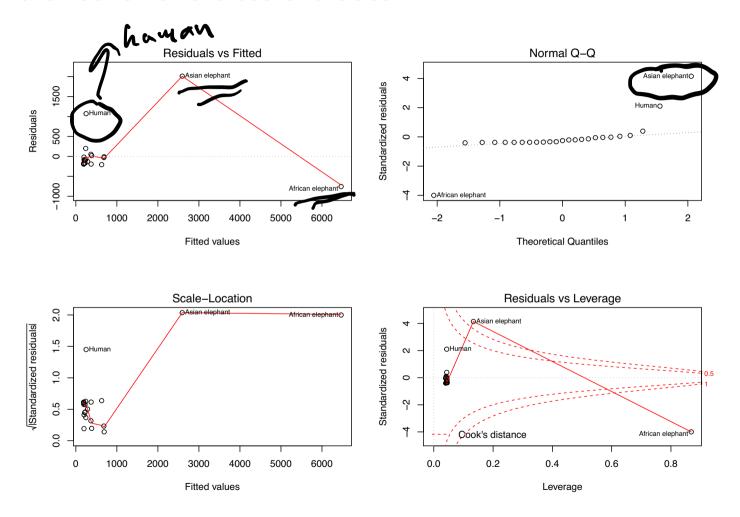
#### Remove Influential Point & Refit



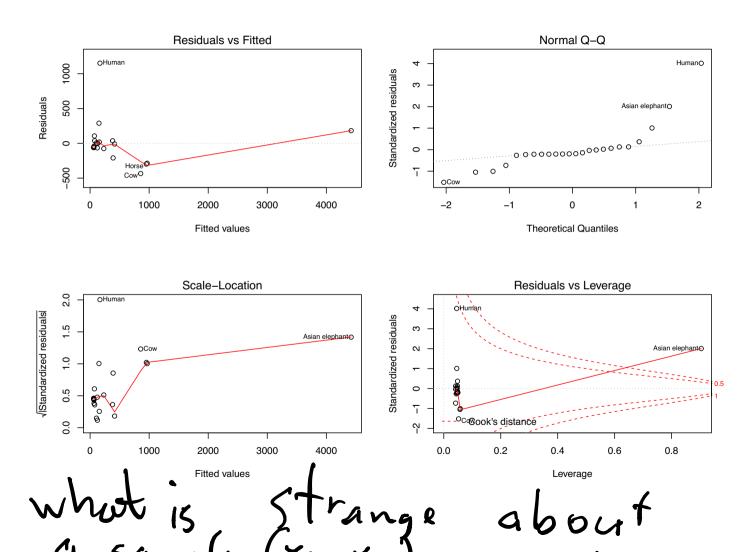
# Keep removing points?



### And another one bites the dust



### and another one



And they just keep coming!

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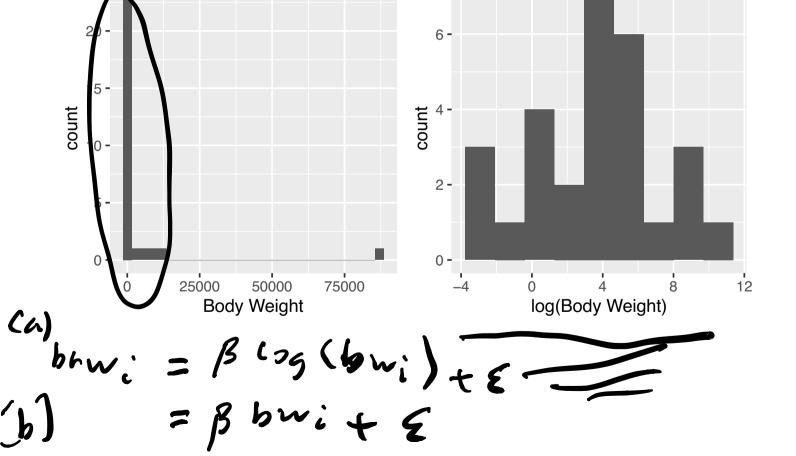
And they just keep coming!



# Plot of Original Data (what you should always do first!)

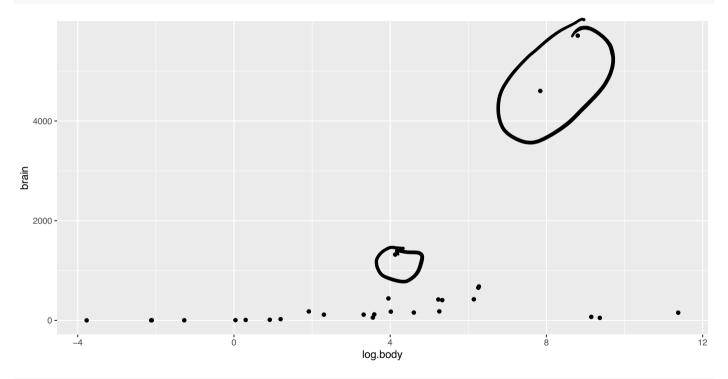
```
library(ggplot2)
ggplot(Animals, aes(x=body, y=brain)) +
  geom_point() +
  xlab("Body Weight (kg)") + ylab("Brain Weight (g)")
                  5 tep-down
                          10× W4/45
 2000 -
                   25000
                            Body Weight (kg)
```

# Log Transform



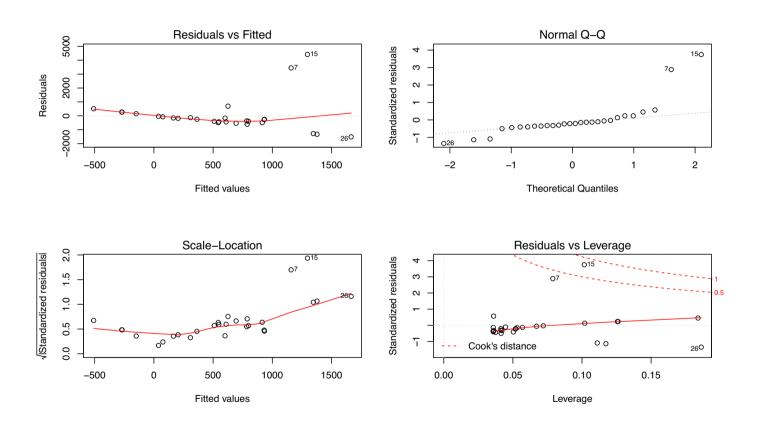
### Plot of Transformed Data

```
Animals= mutate(Animals, log.body = log(body))
ggplot(Animals, aes(log.body, brain)) + geom_point()
```



```
#plot(brain ~ body, Animals, log="x")
```

# Diagnostics with log(body)

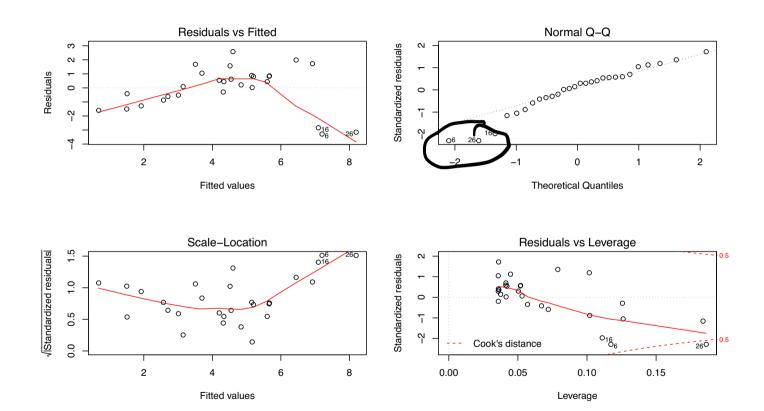


Variance increasing with mean

## Try Log-Log

```
Animals = mutate(Animals, log.brain = log(brain))
ggplot(Animals, aes(log.body, log.brain)) + geom_point()
log.brain
                             e Lepha
#plot(brain ~ body; Anguels, log="xy")
                                             brain is
                                             mulhin
```

# Diagnostics with log(body) & log(brain)



## **Optimal Transformation for Normality**

The BoxCox procedure can be used to find "best" power transformation  $\lambda$  of Y (for positive Y) for a given set of transformed predictors.

$$\Psi(\mathbf{Y}, \lambda) = \begin{cases} \frac{\mathbf{Y}^{\lambda} - 1}{\lambda} & \text{if } \lambda \neq 0 \\ \log(\mathbf{Y}) & \text{if } \lambda = 0 \end{cases}$$

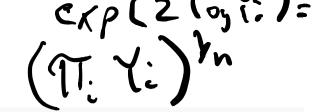
Find value of  $\lambda$  that maximizes the likelihood derived from  $\Psi(\mathbf{Y},\lambda) \sim \mathsf{N}(\mathbf{X}\boldsymbol{\beta}_{\lambda},\sigma_{\lambda}^2)$  (need to obtain distribution of  $\mathbf{Y}$  first)

Find  $\lambda$  to minimize  $|\mathbf{X}\boldsymbol{\beta}_{\lambda}, \boldsymbol{\delta}_{\lambda}| \text{ (Need to obtain distribution of Y first)}$   $|\mathbf{RSS}(\lambda) = \|\Psi_{M}(\mathbf{Y}, \lambda) - \mathbf{X}\hat{\boldsymbol{\beta}}_{\lambda}\|^{2}$ 

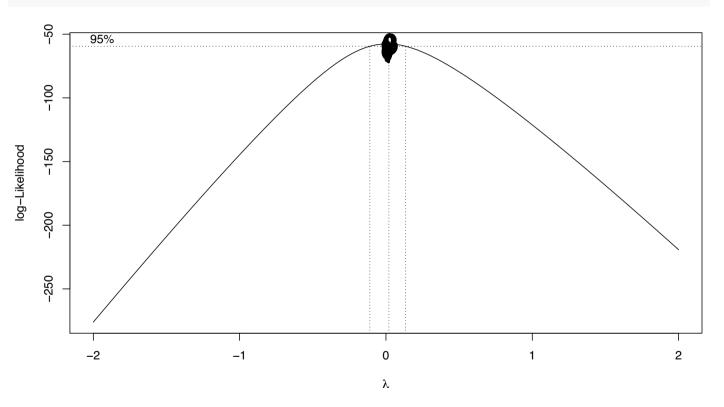
$$\Psi_M(\mathbf{Y},\lambda) = \left\{ egin{array}{ll} (\mathsf{GM}(\mathbf{Y})^{1-\lambda}(\mathbf{Y}^{\lambda}-1)/\lambda & ext{if } \lambda 
eq 0 \ \mathsf{GM}(\mathbf{Y})\log(\mathbf{Y}) & ext{if } \lambda = 0 \end{array} 
ight.$$

where  $GM(\mathbf{Y}) = \exp(\sum \log(Y_i)/n)$  (Geometric mean)

### boxcox in R: Profile likelihood



#### MASS::boxcox(braintransX.lm)



#### Caveats

- ightharpoonup Boxcox transformation depends on choice of transformations of X's
- ► For choice of X transformation use boxTidwell in library(car) or for multivariate normality of response and predictors powerTransform in the car library.
- transformations of X's can reduce leverage values (potential influence)
- if the dynamic range of Y or X is less than 1 or 10 (ie max/min) then transformation may have little effect
- transformations such as logs may still be useful for interpretability

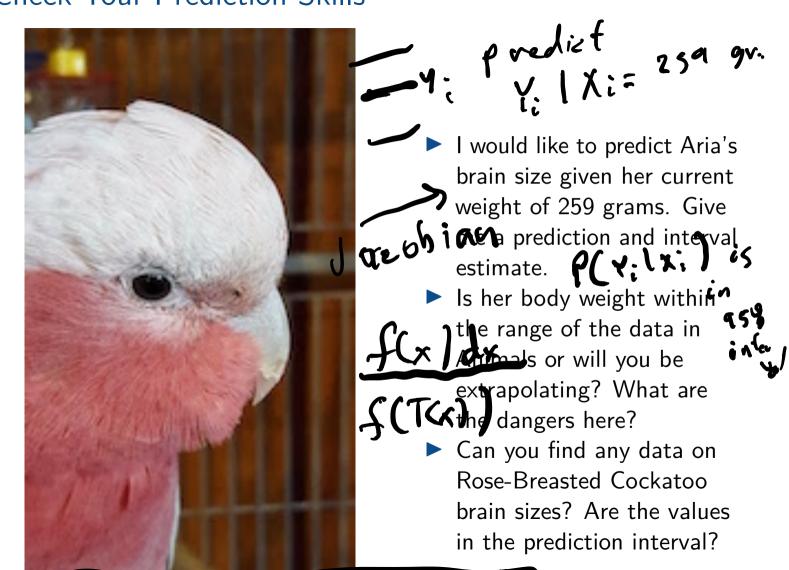


- ightharpoonup outliers that are not influential may still affect the estimate of  $\sigma$  and width of confidence/prediction intervals.
- Reproducibility describe steps and decisions for removing a case

#### Next Class

- ► In the model with both response and predictor log transformed, are dinosaurs outliers?
- should you test each one individually or as a group; if as a group how do you think you would you do this using Im?
- do you think your final model is adequate? What else might you change?
- after you determine whether dinos can stay or go and refine your model, what about prediction?
- what about model uncertainty?

### Check Your Prediction Skills



extrapolation vs. interpolation

modeling

mammals + dinosaurs

log Y: = (log xi) βm + E

log Y: = (log xi) βs + E

don't Know if (xi, yi) is

a dino or a mammal?

hicrarchical model.

Zi=1if dino 0 = wi

Zi-Bern(p)

Yilzi:1, Pd, Xi, oz ~ N(Pd log(xi), oz)

Yilzi:0, Pm, Xi, oz ~ N(Pm (og(xi), oz))

what it I don't give you Zi's? Max (og lik(Y: 1xi, ls, Bm, 0°, Zi) Max B's, Zi ( harder problem