

Analyzing Data With Regression Models

Professor Diane Lambert

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Structure of This Course

The Professor (me)

Past:

Associate Professor of statistics at CMU

worked with environmental safety data

Research Scientist, Bell Labs

network data, semiconductor data

Now:

Research Scientist at Google in New York

network data, data on advertisements, advertisers, image search, and more

The Teaching Assistants

Xiaoxing Cheng (Daniel)

Tianxi Li

Jianghan Qu (Jenny)

Bihyuan Zhang (Elise)

Zoey Zhao

The teaching assistants will help you with the labs.

The Students?

Please send an email to

`pkusms2010@gmail.com`

With

your name

your university

statistics courses you've taken

what days you have exams this week

have you used R before? a little? a lot?

What This Course Is About

Data analysis with regression models

Linear regression

Logistic regression

Multilevel regression

what all these models are

how to work with them in R

how to interpret them

Organization of the Course

Lectures

I will present the models, with examples

You can ask questions during class

Slides and R code used in the lectures will be available after class

Organization of the Course (continued)

Labs

A chance to analyze data in R using the models from lecture.

Assignments will be done in teams of 2 or 3 students.

The teaching assistants and I will help you.

The TAs will explain how to hand in assignments.

Only English will be used in the course.

This is a good opportunity to practice for visiting the US or studying in the US.

Schedule

Lectures 10:00 - 12:00 (with a break)

Labs 14:00 - 16:00

Tentative Plan

		10-12	2-4
June 15, 16	linear regression	✓	✓
June 17	logistic regression	✓	
June 18	logistic regression		✓
June 21	Google statistics		✓
June 22	logistic regression	✓	✓
June 23, 24	multilevel models	✓	✓

Part 1

A regression model answers a question about data.

Basic Parts of A Regression Problem

1. An **outcome** Y that varies
2. Variables X_1, X_2, \dots, X_k that affect Y
These are conditions that may affect Y
Often called **predictors** or **model terms**
3. A **question** about Y
what is the average Y under different conditions
how do different conditions affect Y
how to predict Y under different conditions

More simply

A regression problem tries to answer a question about an outcome Y with data on Y and $\mathbf{X} = (X_1, \dots, X_K)$.

An Example Question

What affects the sleep a mammal needs?

Y : sleep (mean hours/day for a species)

X_1 : body (mean body weight in kg)

X_2 : brain (mean brain weight in g)

X_3 : life (max lifetime in years)

X_4 : predation (1 = low, 5 = high)

X_5 : gestation (days until birth)

fact: Real data analysts give variables real names.

X_1, \dots, X_k are not real names.

Example Data

	sleep	body	brain	predation	danger
African_elephant	3.3	6654.0	5712.0	3	3
African_giant_pouched_rat	8.3	1.0	6.6	3	3
Arctic_Fox	12.5	3.4	44.5	1	1
Asian_elephant	3.9	2547.0	4603.0	3	4
Baboon	9.8	10.6	179.5	4	4
Big_brown_bat	19.7	0.0	0.3	1	1
Brazilian_tapir	6.2	160.0	169.0	4	4
Cat	14.5	3.3	25.6	1	1
Chimpanzee	9.7	52.2	440.0	1	1
Chinchilla	12.5	0.4	6.4	5	4
Cow	3.9	465.0	423.0	5	5
Donkey	3.1	187.1	419.0	5	5
Eastern_American_mole	8.4	0.1	1.2	1	1
Echidna	8.6	3.0	25.0	2	2

There are 51 species in the data (most not shown here).

A Statistical Regression Model

A statistical regression model consists of

1. a model for how the mean outcome Y changes with X_1, \dots, X_K

$$E(Y \mid X_1, \dots, X_K) = \mu(X_1, \dots, X_K)$$

2. a model for how the variability in Y changes with X_1, \dots, X_K

$$\text{var}(Y \mid X_1, \dots, X_K) = \sigma^2(X_1, \dots, X_K)$$

A regression model may or may not assume

3. a distribution for $Y \mid X_1, \dots, X_K$.

Simplest Regression Model For Sleep

Mean

$$E(\text{sleep} \mid \mathbf{X}) = \mu$$

Variance

$$\text{var}(\text{sleep} \mid \mathbf{X}) = \sigma^2$$

Distribution

$$\text{sleep} \sim \text{Normal}(\mu, \sigma^2)$$

X isn't used to predict sleep in this model

Is this a good enough model for sleep
across species of mammals?

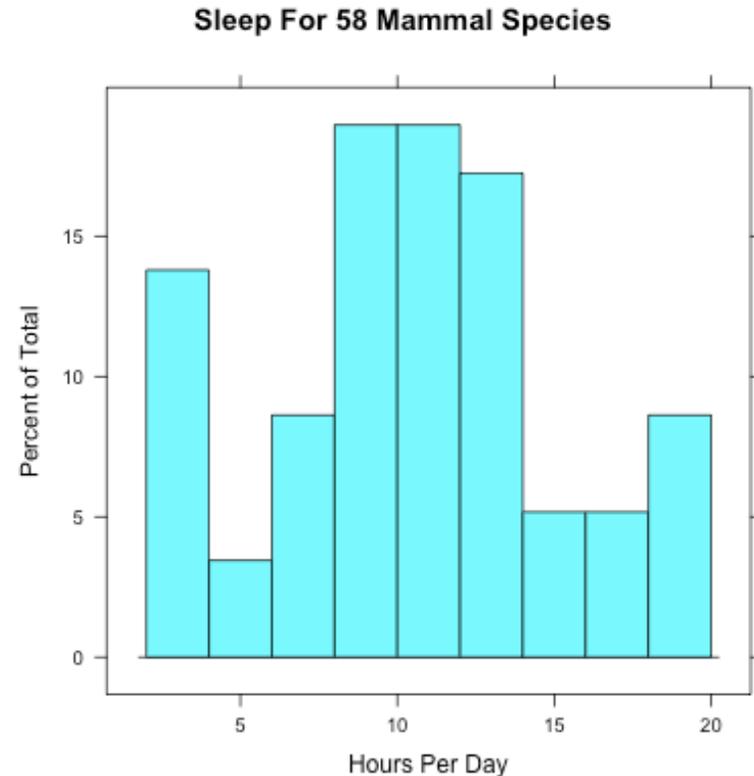
The Simplest Model For Sleep

$$E(\text{sleep} \mid \mathbf{X}) = \mu$$

$$\text{var}(\text{sleep} \mid \mathbf{X}) = \sigma^2$$

$$\text{sleep} \sim \text{Normal}(\mu, \sigma^2)$$

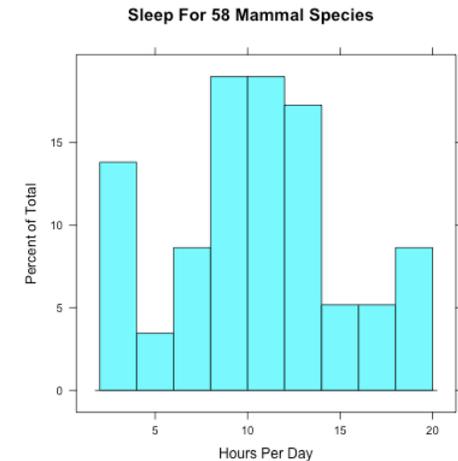
If the model is reasonable, the sleep data should look like a random sample from a normal distribution



The histogram doesn't look like a normal curve, so the simplest model isn't very good.

To Get Better Models

We could use a different distribution
we need more mass in the tails
it is hard to find a distribution like
that, and it won't give us much
insight into why there is mass in the
tails.



Instead, try to model the mean
we can use the other information
(like body weight) about the species

Making The Mean Depend on \mathbf{X}

Linear regression has two assumptions.

1. Mean outcome Y is linear in the terms in \mathbf{X}

$$\mu(\mathbf{X}) = \beta_0 + \beta_1 X_1 + \dots + \beta_K X_K$$

2. Variance of the outcome is constant in \mathbf{X} .

$$\text{var}(Y | \mathbf{X}) = \sigma^2$$

Usually, the distribution of $Y|\mathbf{X}$ is unimportant, as long as Y is numeric and has several possible values.

$Y|\mathbf{X}$ does not have to be normally distributed.

Back to The Example

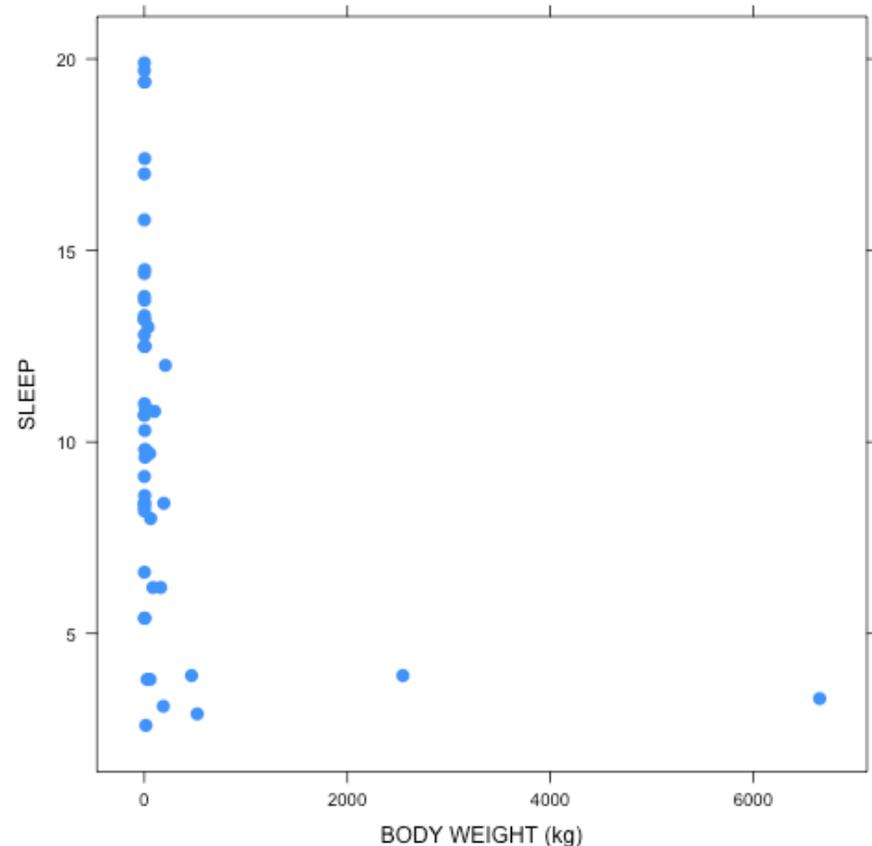
Is the mean sleep of a mammal species linear in body weight?

There is no theory here.

Look at the data. →

Answer:

A line doesn't describe how $E(\text{sleep} \mid \text{body})$ depends on body weight.



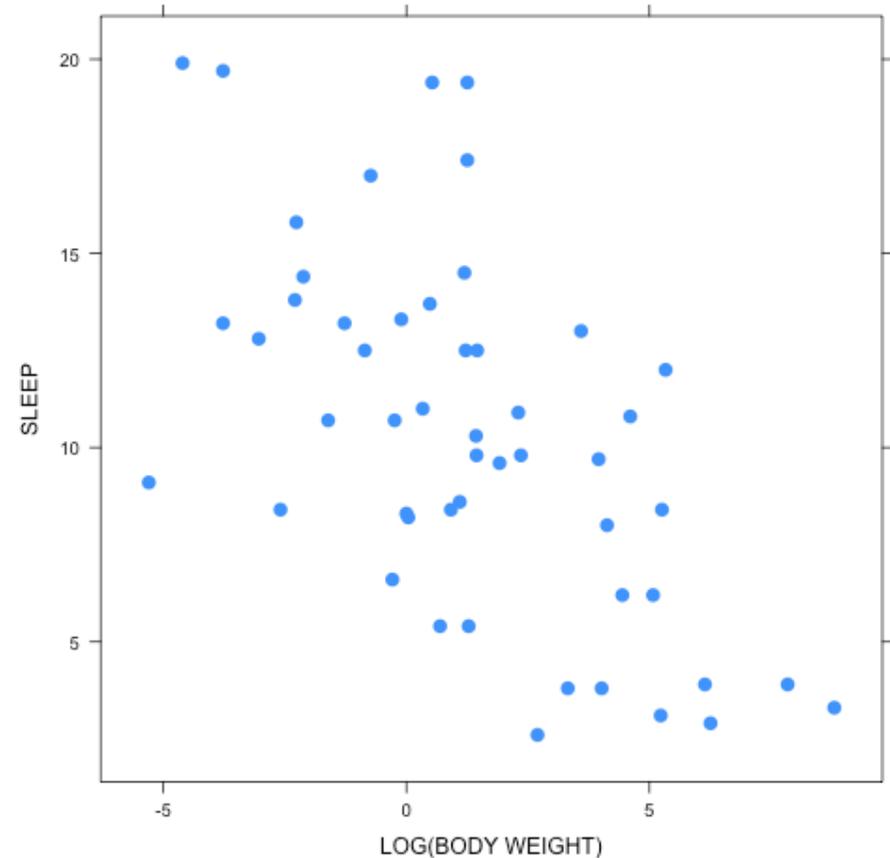
Linear regression of sleep on body weight is not a good model for mammals *even though b_1 is statistically significant here*. Looking at the regression coefficients won't tell you if the model fits.

Changing Predictors

In linear regression, the mean is linear in the predictors.

An X can be transformed, or several can be combined like X_1X_2 and it's still linear regression.

Body weight covers several orders of magnitude. In such cases, try $\log(X)$.



Linear regression of sleep on $\log(\text{body weight})$ is sensible.

Part 2

Linear Regression Coefficients

Least Squares Coefficient Estimates

With **estimated coefficients** b_0, \dots, b_K , the **residuals** or **prediction errors** are

$$e_i = Y_i - b_0 - b_1X_1 - \dots - b_kX_K$$

The **least squares coefficients** b_0, \dots, b_K minimize the sum of squared errors

$$\sum_i (Y_i - b_0 - b_1X_1 - \dots - b_kX_K)^2$$

FACT: The b_k 's are also maximum likelihood estimates when $Y|(X_0, \dots, X_K)$ has a normal distribution. In practice, estimates are often chosen to minimize a loss function like mean squared error without assigning a distribution to Y .

Linear Regression Without Predictors

The estimated intercept

$$b_0 = \text{sample mean}$$

The residuals

$$e_i = Y_i - b_0$$

The estimated $\text{var}(Y | X)$ [X is null]

$$s^2 = (n - 1)^{-1} \sum_{i=1}^n (y_i - b_0)^2$$

For the sleep data,

$$b_0 = 10.3, \quad s^2 = 21.9$$

Computing Regression Coefficients

Any reasonable statistical software will compute the least squares coefficients (b 's) and residuals (e 's).

We'll use R -- it's commonly used in research

Many corporations use SAS -- I don't know SAS.

Fact (don't worry if you haven't seen it before):

$$\mathbf{b} = (\mathbf{X}^t \mathbf{X})^{-1} \mathbf{X}^t \mathbf{Y}$$

where

\mathbf{b} is the vector of K estimated coefficients

\mathbf{Y} is the vector of n outcomes (one for each observation)

\mathbf{X} is an $n \times K$ matrix, each row is a different observation and each column is a different predictor

Estimates For The Sleep Data

The estimated line

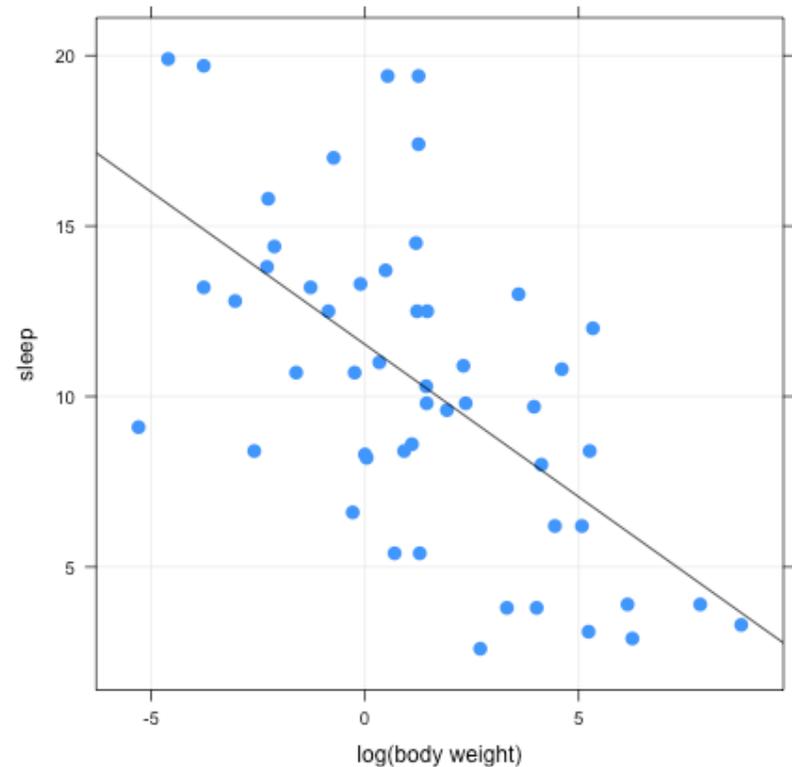
$$\mu(X_1) = b_0 + b_1 X_1$$

has coefficients

$$(b_0, b_1) = (11.5, -0.9)$$

a species with mean weight 1 kg is estimated to sleep 11.5 hours on average

a species with a mean weight of 100 kg is estimated to sleep 7.4 hours on average.



Residuals for the Sleep Data

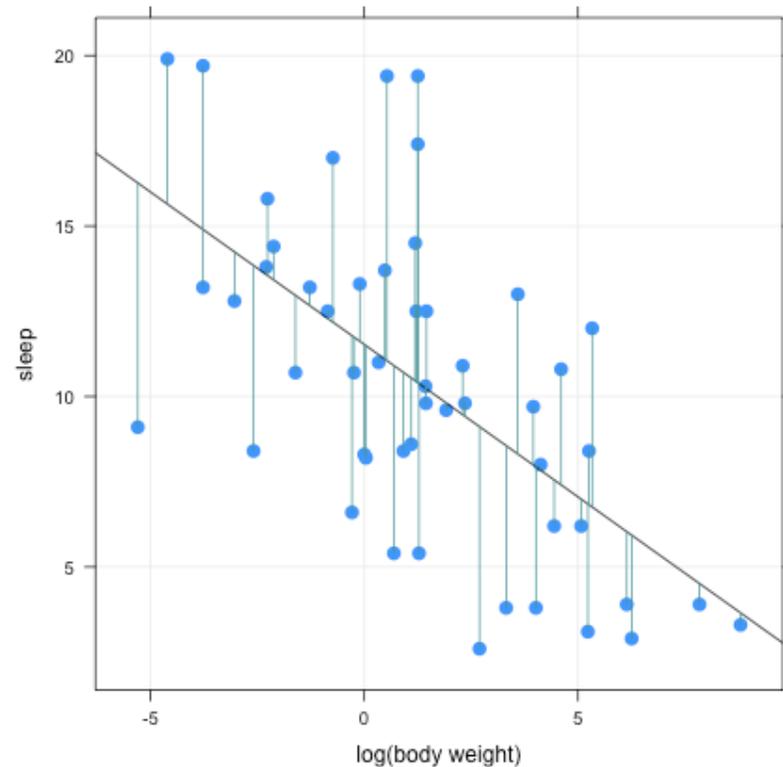
Gray segments show
the residuals

$$e_i = Y_i - b_0 - b_1 \log(\text{body}_i)$$

If a segment is below the line, the outcome Y for that species is smaller than its estimated mean.

The closer the residuals are to zero, the better the regression fits the data.

Most measures of how well a model fits the data are based on residuals.



What About $\text{var}(Y | \mathbf{X})$?

In linear regression, the variance of sleep should not depend on \mathbf{X}

$$\text{var}(Y | X_1, \dots, X_k) = s^2$$

Example:

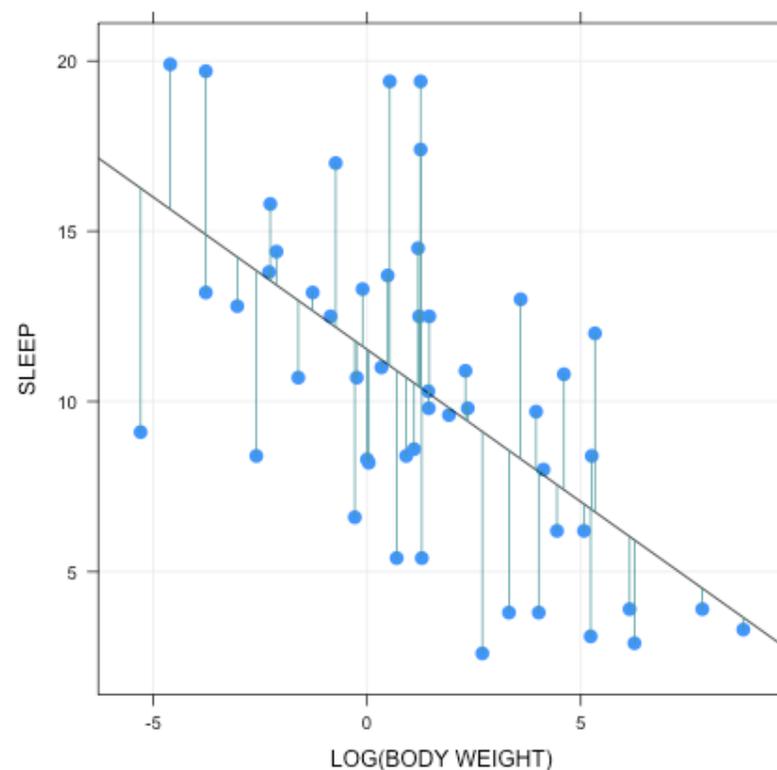
The variance of sleep for species with average weight 100 kg should be the same as the variance of sleep for species with average weight 1 kg.

Is this assumption plausible for the sleep data?

Variance of $Y | X$

If the residuals have a pattern in X , then $\text{var}(Y | X)$ depends on X , so the model is bad.

Here, the residuals don't vary consistently with $\log(\text{body weight})$.



The assumption that variance of sleep is the same for all body weights passes the eyeball test.

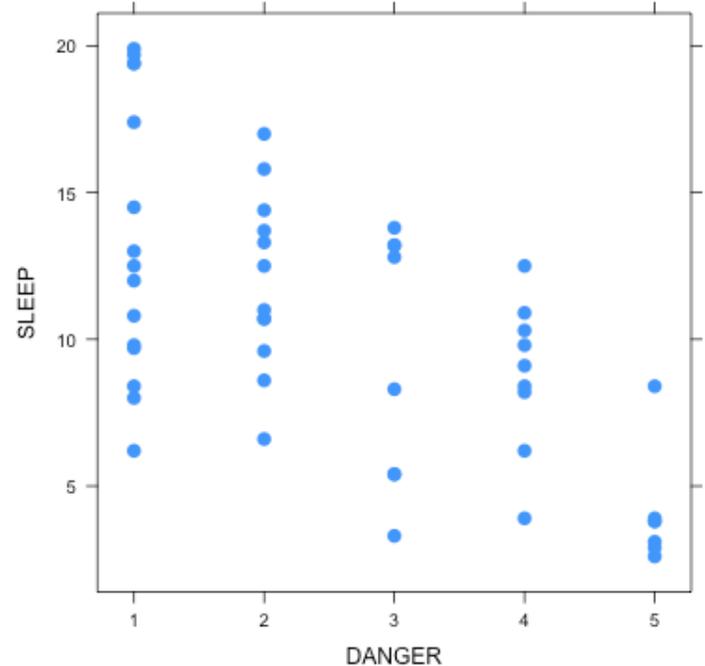
Fact: $\text{var}(Y | X)$ is estimated by $s^2 = (n - K)^{-1} \sum_{i=1}^n e_i^2$

Multiple Predictors

Including more predictors can improve a model

A quick look at the predictor danger →

Fit a regression model with $\log(\text{body})$ and danger as predictors in R.



	intercept	$\log(\text{body})$	danger
old model	11.5	-.89	
new model	15.7	-.67	-1.68

Including a new predictor can change our understanding of an old predictor.

Evaluating the Fit of Linear Regression Models

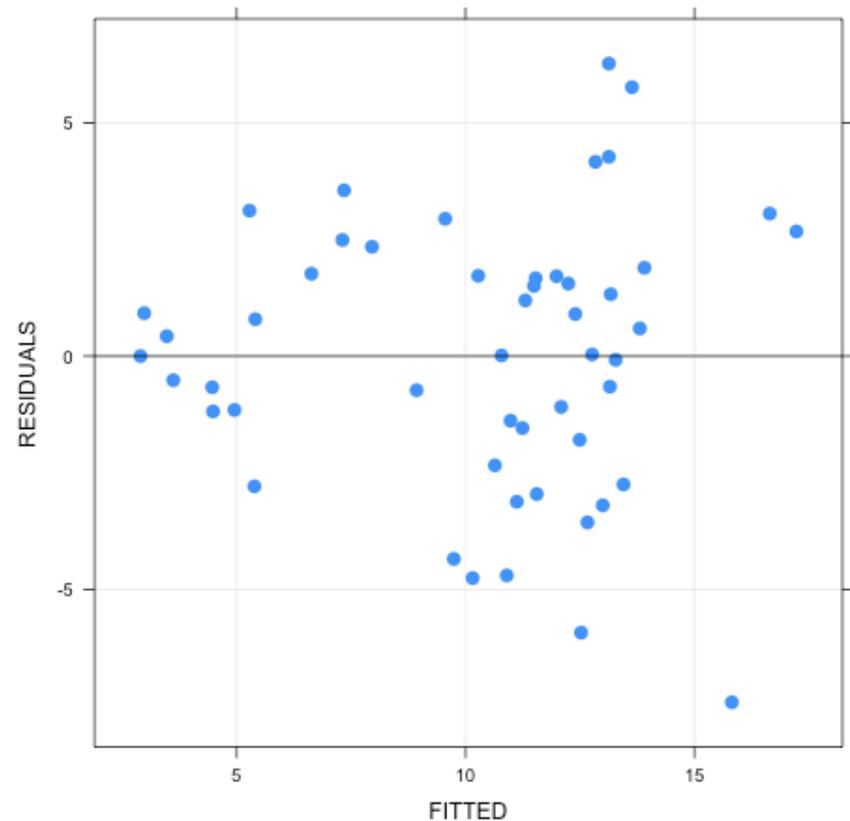
First Look At The Residuals

Residuals vs Estimated Means (the fitted values)

a pattern suggests that the model doesn't fit

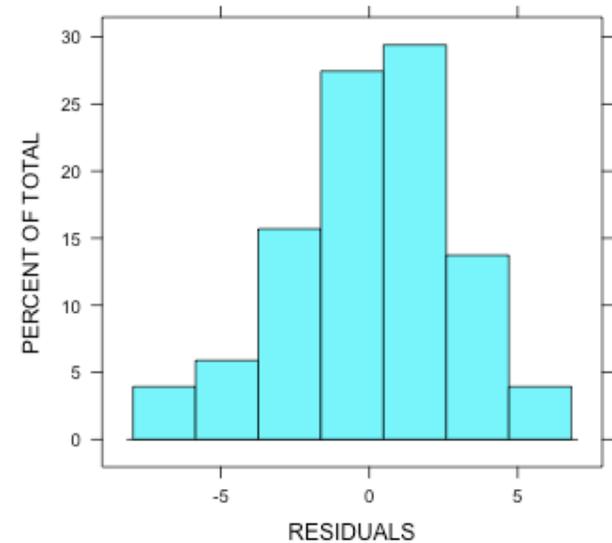
sleep example with $\log(\text{body})$ and danger as predictors

symmetric about zero over the range of the fitted, but a few points suggest taking a closer look at model fit.

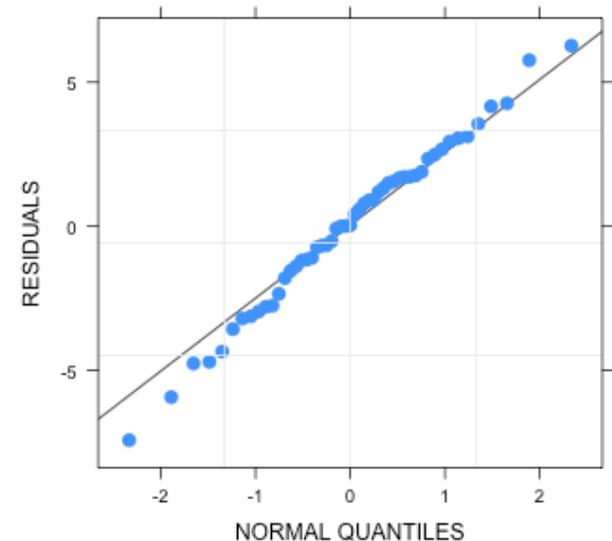


Residuals Should Be Close to Normal

Histogram of residuals looks reasonably similar to that of a random sample from a normal distribution with mean zero.



The ordered residuals look reasonably like the expected order statistics from a normal distribution with mean zero.



Classical Regression Statistics

The linear regression coefficients depend on the data (X_1, \dots, X_k, Y) so have standard errors.

R and SAS compute standard errors.

$b_k/\text{se}(b_k)$ is approximately t_{n-k} when $\beta_0=0$

Warning: The t-statistics assume that the other terms are in the model.

	b_k	$\text{se}(b_k)$	t-value	$\text{Pr}(> t)$
(Intercept)	15.6902	0.8771	17.889	$< 2e-16$
log(body)	-0.6993	0.1368	-5.112	$5.50e-06$
danger	-1.6830	0.3041	-5.534	$1.28e-06$

More Classical Regression Statistics

We can get confidence intervals for the mean Y and the prediction for a different species under conditions X_1, \dots, X_K .

body	danger	estimate	meanLow	meanHigh	predLow	predHigh
0.1	1	15.6	14.1	17.1	9.5	21.8
1.0	1	14.0	12.7	15.3	7.9	20.1
10.0	1	12.4	11.0	13.8	6.3	18.5
100.0	1	10.8	9.1	12.5	4.6	17.0
0.1	5	8.9	6.8	11.0	2.6	15.2
1.0	5	7.3	5.5	9.1	1.0	13.5
10.0	5	5.7	4.0	7.3	-0.5	11.8
100.0	5	4.1	2.3	5.8	-2.2	10.3

The predicted value is the mean, but predicting is riskier than estimating a mean.