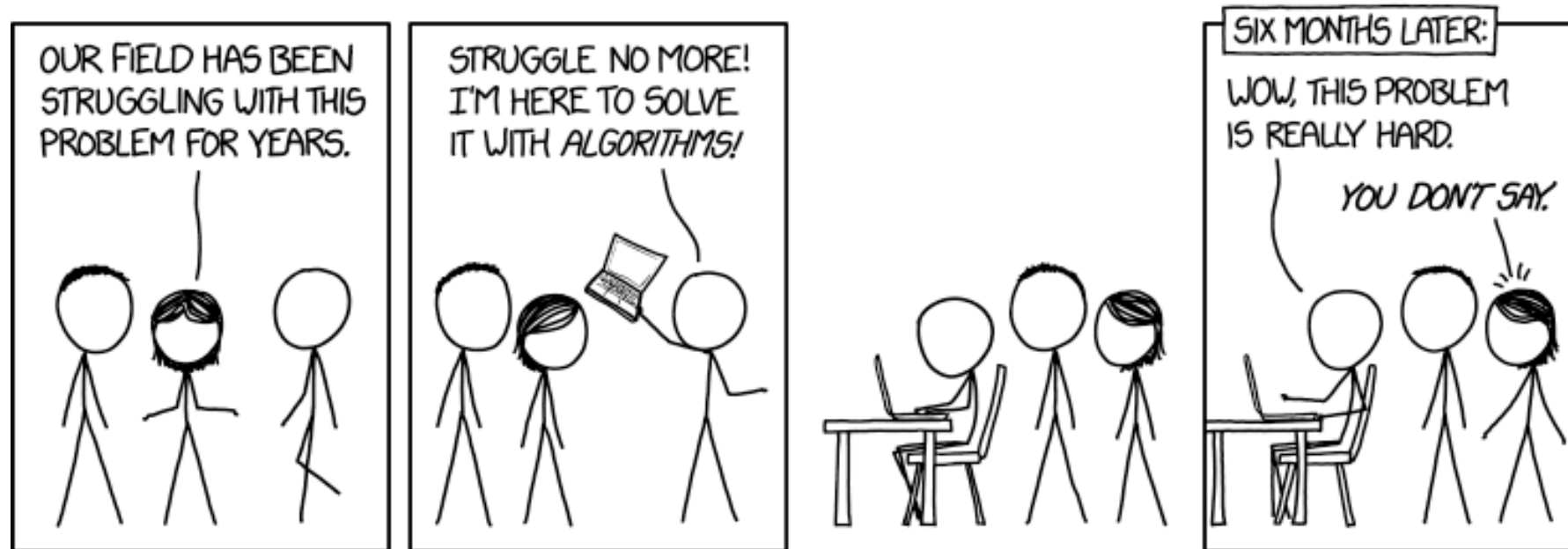


# Multiple Linear Regression

ENTMLGY 6707 Entomological Techniques and Data Analysis



# Learning objectives

- 1) Compare and contrast simple vs multiple linear regression
- 2) Become familiar with additional assumptions when using multiple linear regression and how to check for them / deal with them.
- 3) Interpret the outcome of a multiple linear regression

# Simple linear regression

**Table 4.1** Examples of the Generalized Linear Model as a Function of Independent Variable and Dependant Variable Type

		Responses				
		<i>Continuous DV</i>	<i>Binary DV</i>	<i>Unordered Multicategory DV</i>	<i>Ordered Categorical DV</i>	<i>Count DV</i>
Predictors	<i>Continuous IV</i>	OLS regression	Binary logistic regression	Multinomial logistic regression	Ordinal logistic regression	OLS, Poisson regression
	<i>Mixed continuous and categorical IV</i>					
	<i>Binary/ categorical IV only</i>	ANOVA and <i>t</i> -test	Log-linear models	Log-linear models		Log-linear models

ANOVA, analysis of variance; DV, dependent variable; IV, independent variable; OLS, ordinary least squares.

Chapter 4: Simple Linear Models With Continuous Dependent Variables: Simple ANOVA Analyses

In: [Regression & Linear Modeling: Best Practices and Modern Methods](#)

# Multiple linear regression

**Table 4.1** Examples of the Generalized Linear Model as a Function of Independent Variable and Dependant Variable Type

		Responses				
		<i>Continuous DV</i>	<i>Binary DV</i>	<i>Unordered Multicategory DV</i>	<i>Ordered Categorical DV</i>	<i>Count DV</i>
Predictors	<i>Continuous IV</i>	OLS regression	Binary logistic regression	Multinomial logistic regression	Ordinal logistic regression	OLS, Poisson regression
	<i>Mixed continuous and categorical IV</i>					
	<i>Binary/ categorical IV only</i>	ANOVA and <i>t</i> -test	Log-linear models	Log-linear models		Log-linear models

ANOVA, analysis of variance; DV, dependent variable; IV, independent variable; OLS, ordinary least squares.

Chapter 4: Simple Linear Models With Continuous Dependent Variables: Simple ANOVA Analyses

In: [Regression & Linear Modeling: Best Practices and Modern Methods](#)

# Linear regression

$$Y_i = \beta_0 + \beta_1 X_i + \epsilon_i$$

Response variable

**y-intercept** (or the expected mean of  $y$  when  $x=0$ )

**Slope coefficient** (rise over run, or the expected change in  $y$  with a 1 unit change in  $x$ )

Predictor variable

Residuals

The diagram shows the linear regression equation  $Y_i = \beta_0 + \beta_1 X_i + \epsilon_i$ . Arrows point from descriptive labels to each term in the equation:  $Y_i$  is the response variable,  $\beta_0$  is the y-intercept,  $\beta_1$  is the slope coefficient,  $X_i$  is the predictor variable, and  $\epsilon_i$  represents the residuals.

# Comparing simple vs. polynomial linear regression

Simple linear regression

$$Y_i = \beta_0 + \beta_1 X_1 + \varepsilon_i$$

$$\textit{Height} \sim \textit{DBH}$$

Polynomial regression

$$Y_i = \beta_0 + \beta_1 X_{\textcolor{red}{1}} + \beta_2 X_{\textcolor{red}{1}}^2 + \varepsilon_i$$

$$\textit{Height} \sim \textit{DBH} + \textit{DBH}^2$$

# Comparing polynomial vs. multiple linear regression

Polynomial regression

$$Y_i = \beta_0 + \beta_1 X_{\textcolor{red}{1}} + \beta_2 X_{\textcolor{red}{1}}^2 + \varepsilon_i$$

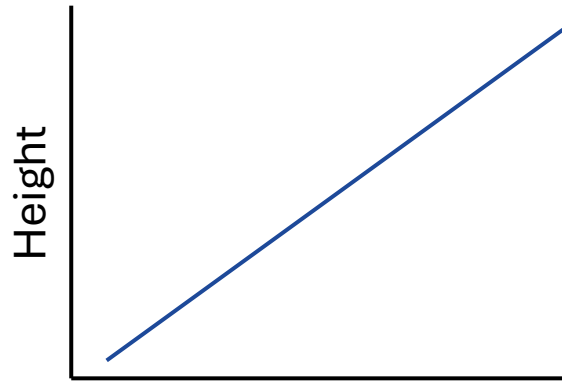
$$\textit{Height} \sim \textit{DBH} + \textit{DBH}^2$$

Multiple linear regression

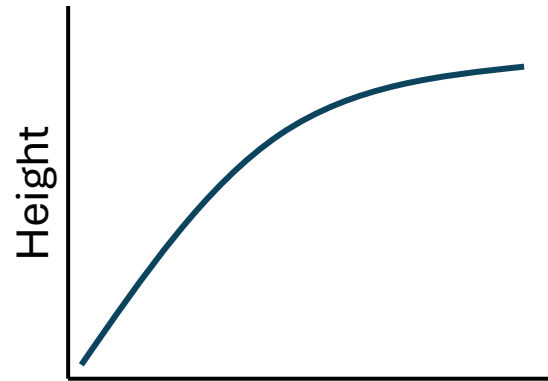
$$Y_i = \beta_0 + \beta_1 X_{\textcolor{red}{1}} + \beta_2 X_{\textcolor{red}{2}} + \varepsilon_i$$

$$\textit{Height} \sim \textit{DBH} + \textit{Nitrogen}$$

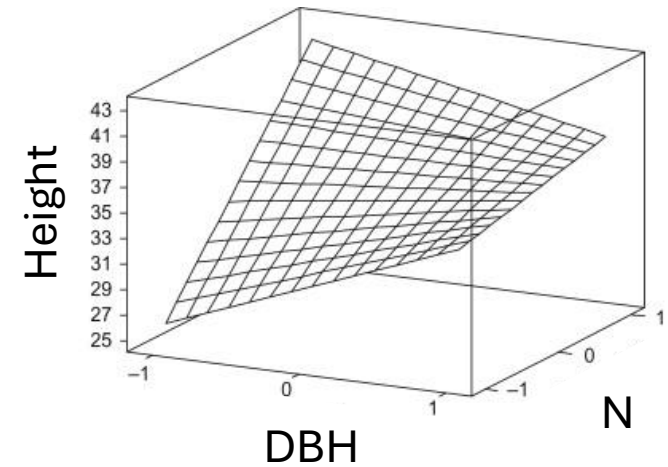
# Single vs. polynomial vs. multiple regression models



DBH  
 $Height \sim DBH$



DBH  
 $Height \sim DBH + DBH^2$



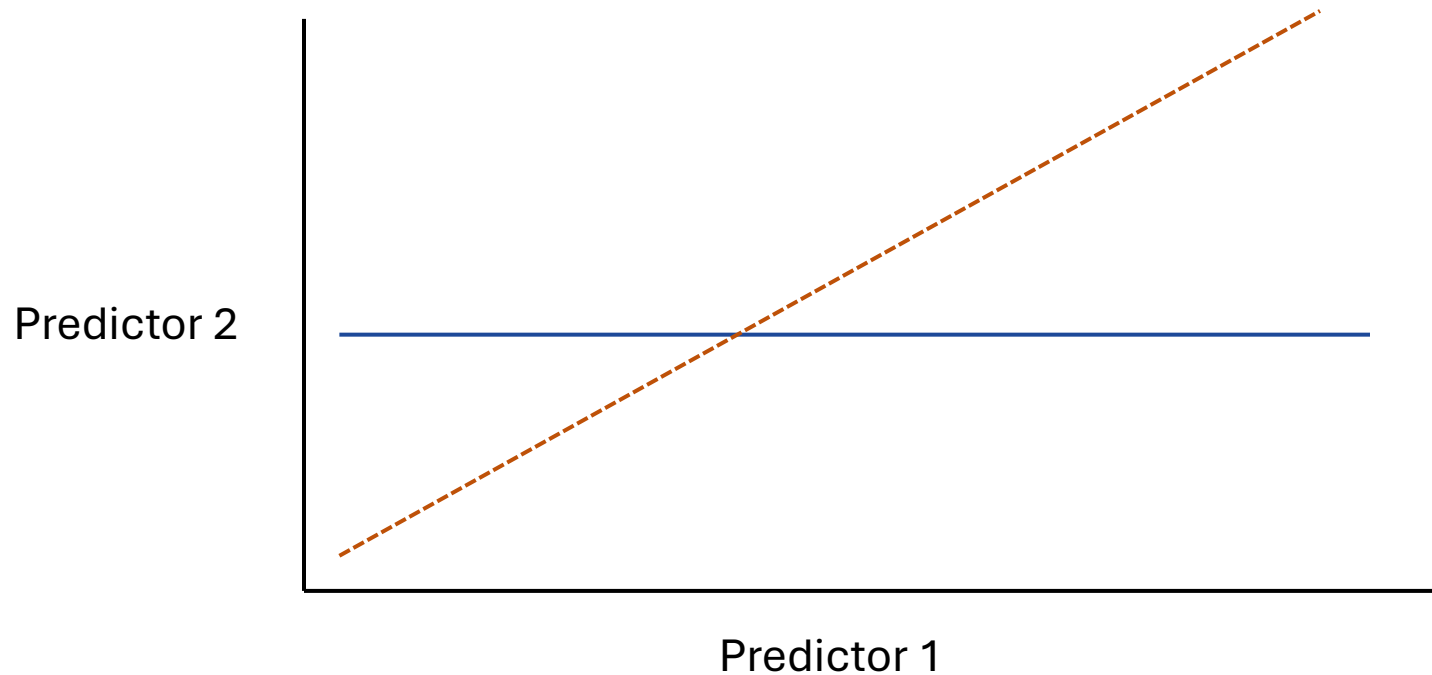
DBH N  
 $Height \sim DBH + Nitrogen$



# Multiple regression models: additional assumption

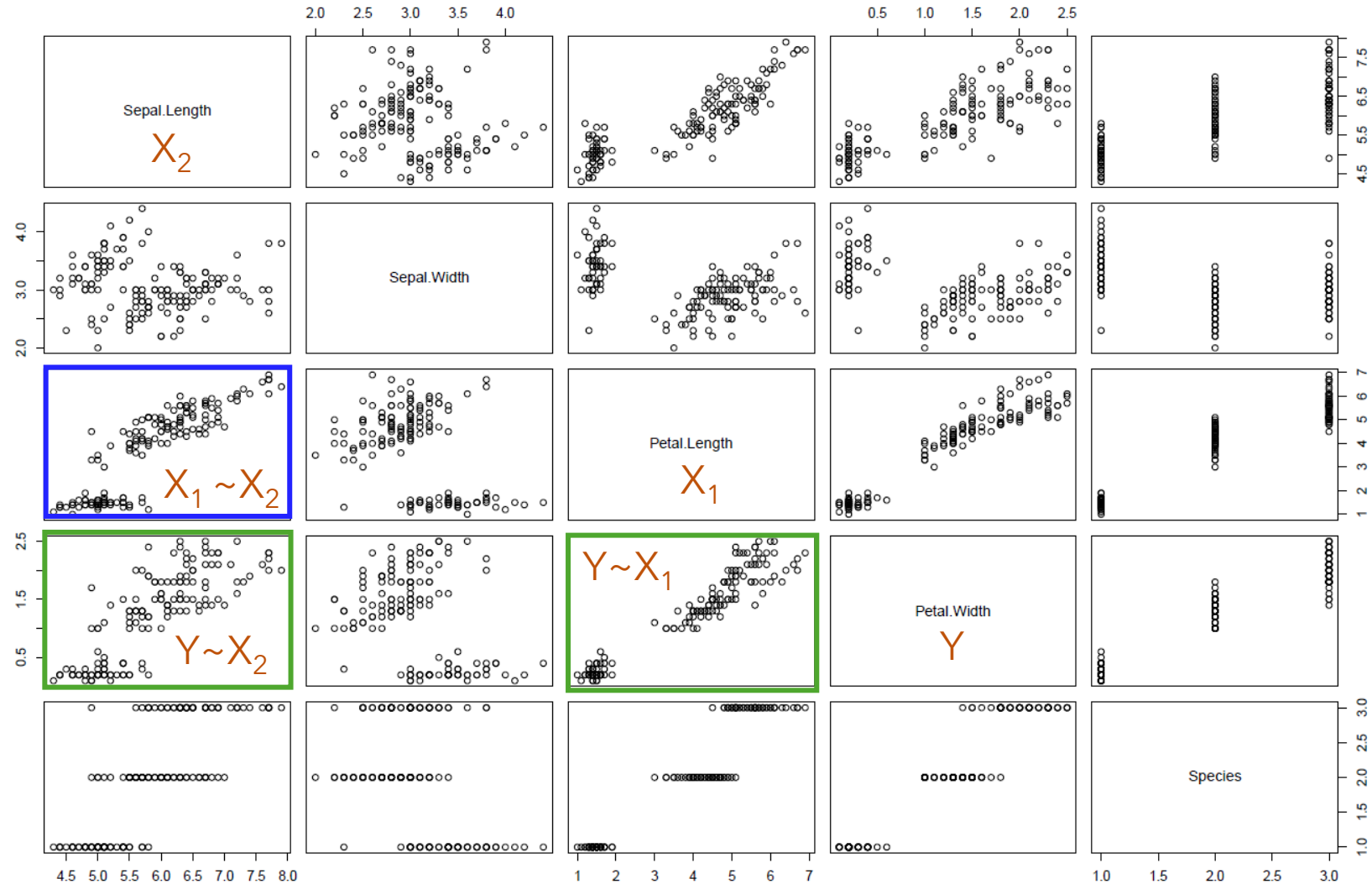
A correlation – negative or positive – between predictors is called **collinearity** which can cause problems in model fitting.

A common sign of collinearity is “large” changes in slope coefficients, including sign flipping (e.g., slope coefficient goes from negative to positive), depending on which predictors are fit in a model.



$$Petal.Width \sim Petal.Length + Sepal.Length$$

```
plot(iris)
```



```
fitA <- lm(Petal.Width ~ Petal.Length, data = iris)
summary(fitA)
```

Call:

```
lm(formula = Petal.Width ~ Petal.Length, data = iris)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.56515	-0.12358	-0.01898	0.13288	0.64272

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-0.363076	0.039762	-9.131	4.7e-16	***
Petal.Length	0.415755	0.009582	43.387	< 2e-16	***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.2065 on 148 degrees of freedom

Multiple R-squared: 0.9271, Adjusted R-squared: 0.9266

F-statistic: 1882 on 1 and 148 DF, p-value: < 2.2e-16

```
fitB <- lm(Petal.Width ~ Sepal.Length, data = iris)
summary(fitB)
```

Call:

```
lm(formula = Petal.Width ~ Sepal.Length, data = iris)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.96671	-0.35936	-0.01787	0.28388	1.23329

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-3.20022	0.25689	-12.46	<2e-16 ***
Sepal.Length	0.75292	0.04353	17.30	<2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.44 on 148 degrees of freedom

Multiple R-squared: 0.669, Adjusted R-squared: 0.6668

F-statistic: 299.2 on 1 and 148 DF, p-value: < 2.2e-16

```
fitC <- lm(Petal.Width ~ Petal.Length + Sepal.Length, data = iris)
summary(fitC)
```

Call:

```
lm(formula = Petal.Width ~ Petal.Length + Sepal.Length, data = iris)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.60598	-0.12560	-0.02049	0.11616	0.59404

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-0.008996	0.182097	-0.049	0.9607	
Petal.Length	0.449376	0.019365	23.205	<2e-16	***
Sepal.Length	-0.082218	0.041283	-1.992	0.0483	*

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.2044 on 147 degrees of freedom

Multiple R-squared: 0.929, Adjusted R-squared: 0.9281

F-statistic: 962.1 on 2 and 147 DF, p-value: < 2.2e-16

```
fitA <- lm(Petal.Width ~ Petal.Length, data = iris)
```

**Coefficients: fitA**

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-0.363076	0.039762	-9.131	4.7e-16	***
Petal.Length	0.415755	0.009582	43.387	< 2e-16	***

```
fitB <- lm(Petal.Width ~ Sepal.Length, data = iris)
```

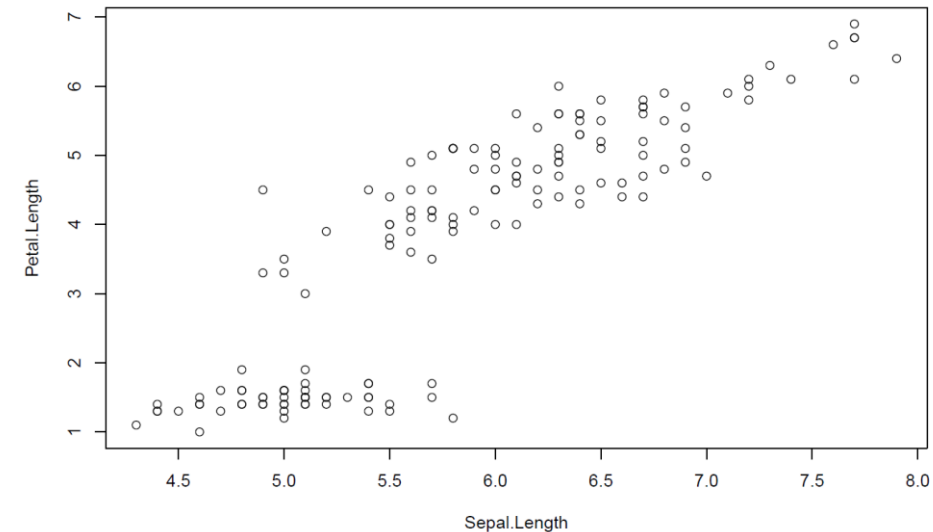
**Coefficients: fitB**

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-3.20022	0.25689	-12.46	<2e-16	***
Sepal.Length	0.75292	0.04353	17.30	<2e-16	***

```
fitC <- lm(Petal.Width ~ Petal.Length + Sepal.Length, data = iris)
```

**Coefficients: fitC**

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-0.008996	0.182097	-0.049	0.9607	
Petal.Length	0.449376	0.019365	23.205	<2e-16	***
Sepal.Length	-0.082218	0.041283	-1.992	0.0483	*



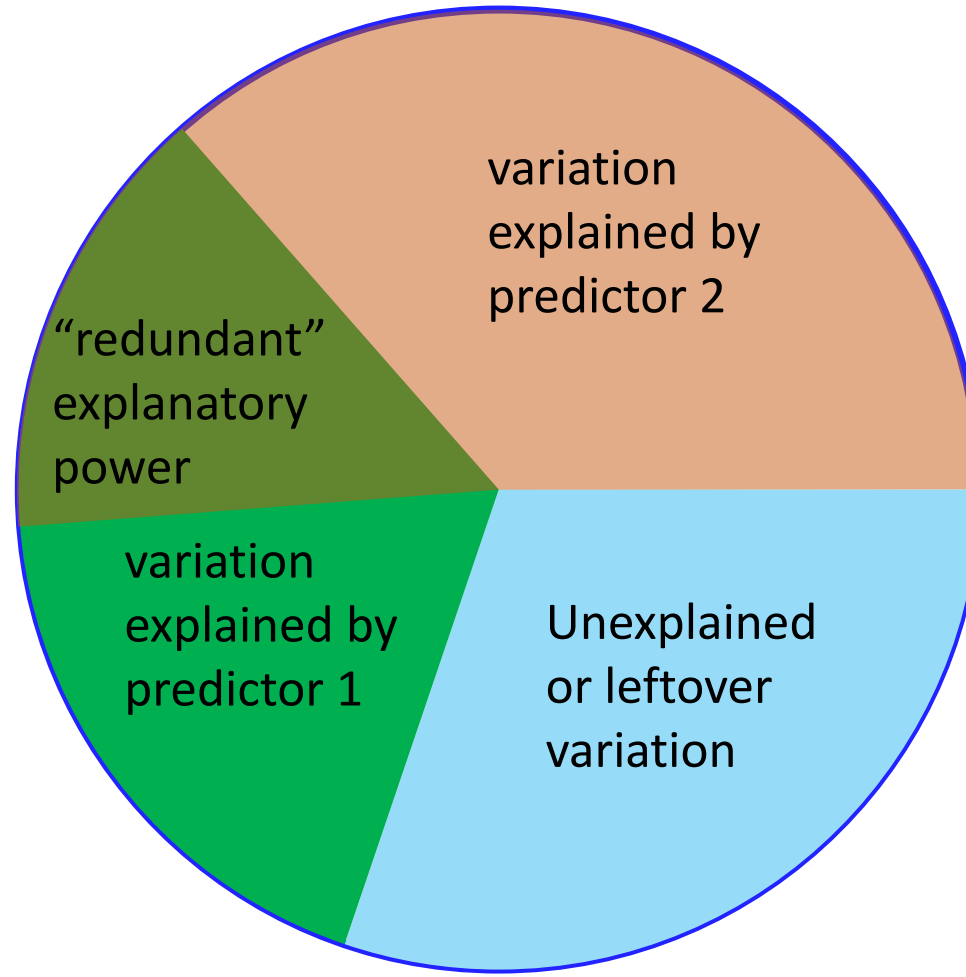
# Interpreting simple vs multiple regression models

In **SLR**, we interpret the slope coefficient as follows: “a one unit increase in  $X_1$  was associated with a  $\beta_1$  unit increase in  $Y$  .”

In **MLR**, we interpret coefficients as follows: “holding all else equal, a one unit increase in  $X_1$  was associated with a  $\beta_1$  unit increase in  $Y$  .”

- That is part of the reason collinearity causes problems. If predictors  $X_1$  and  $X_2$  are highly correlated, it is difficult to “hold  $X_2$  equal or constant” while estimating the effect of  $X_1$ .

# Sequential vs. marginal fits in ANOVA





# Sequential fitting

```
grouseticks$f_YEAR <- as.factor(grouseticks$YEAR)
fit_ex1 <- lm(TICKS ~ f_YEAR + HEIGHT, data=grouseticks)
anova(fit_ex1)
```

```
## Analysis of Variance Table
##
## Response: TICKS
##              Df Sum Sq Mean Sq F value    Pr(>F)
## f_YEAR         2   7050   3524.9   24.995 5.928e-11 ***
## HEIGHT         1   6092   6092.0   43.199 1.550e-10 ***
## Residuals    399  56268    141.0
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
fit_ex2 <- lm(TICKS ~ HEIGHT + f_YEAR, data=grouseticks)
anova(fit_ex2)
```

```
## Analysis of Variance Table
##
## Response: TICKS
##              Df Sum Sq Mean Sq F value    Pr(>F)
## HEIGHT         1   7692   7692.2   54.546 8.948e-13 ***
## f_YEAR         2   5450   2724.8   19.321 9.788e-09 ***
## Residuals    399  56268    141.0
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# Marginal fitting

```
library(car)
```

```
Anova(fit_ex1, type="III")
```

```
## Anova Table (Type III tests)
```

```
##
```

```
## Response: TICKS
```

##	Sum Sq	Df	F value	Pr(>F)	
## (Intercept)	7444	1	52.786	1.970e-12	***
## f_YEAR	5450	2	19.321	9.788e-09	***
## HEIGHT	6092	1	43.199	1.550e-10	***
## Residuals	56268	399			

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Anova(fit_ex2, type="III")
```

```
## Anova Table (Type III tests)
```

```
##
```

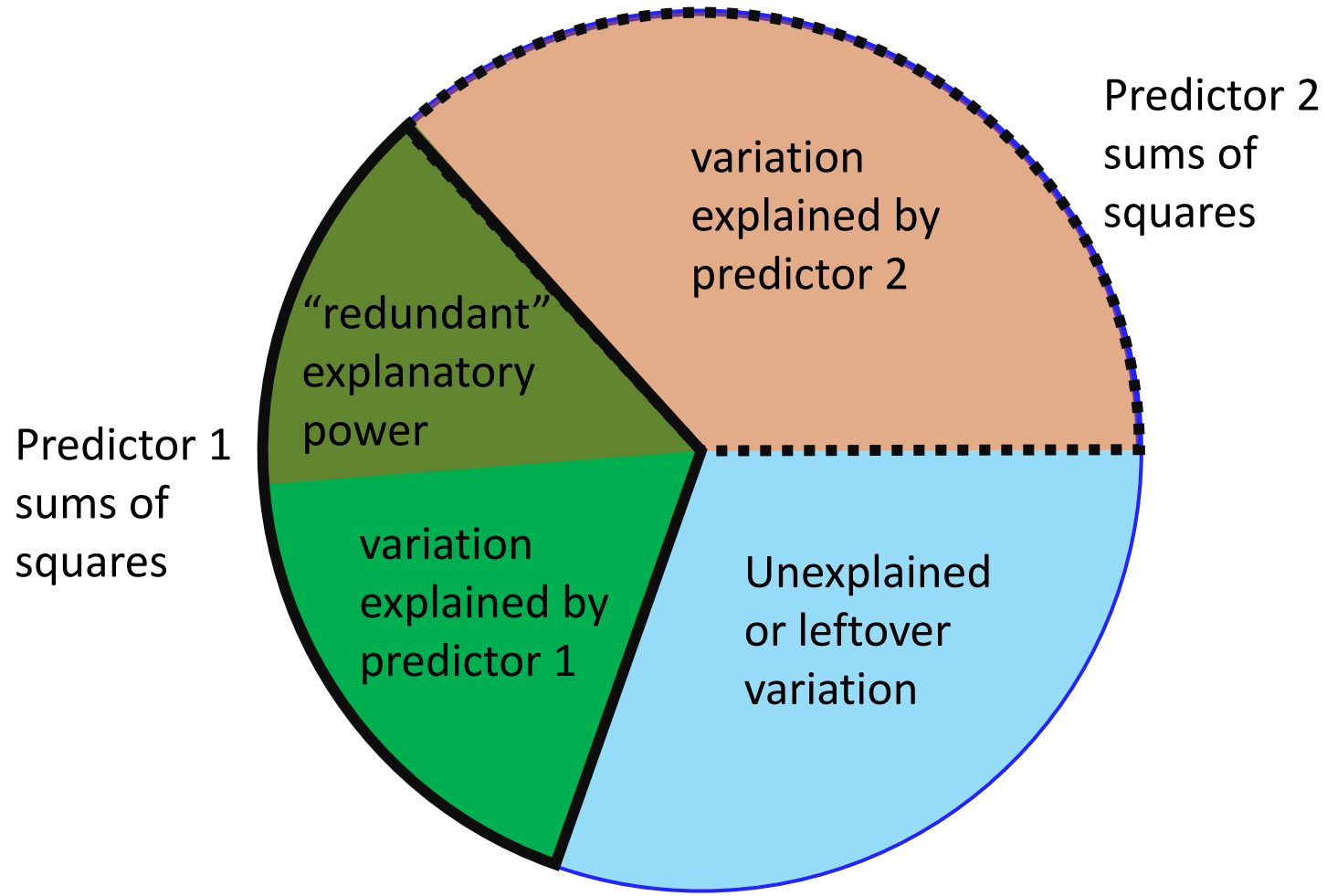
```
## Response: TICKS
```

##	Sum Sq	Df	F value	Pr(>F)	
## (Intercept)	7444	1	52.786	1.970e-12	***
## HEIGHT	6092	1	43.199	1.550e-10	***
## f_YEAR	5450	2	19.321	9.788e-09	***
## Residuals	56268	399			

```
## ---
```

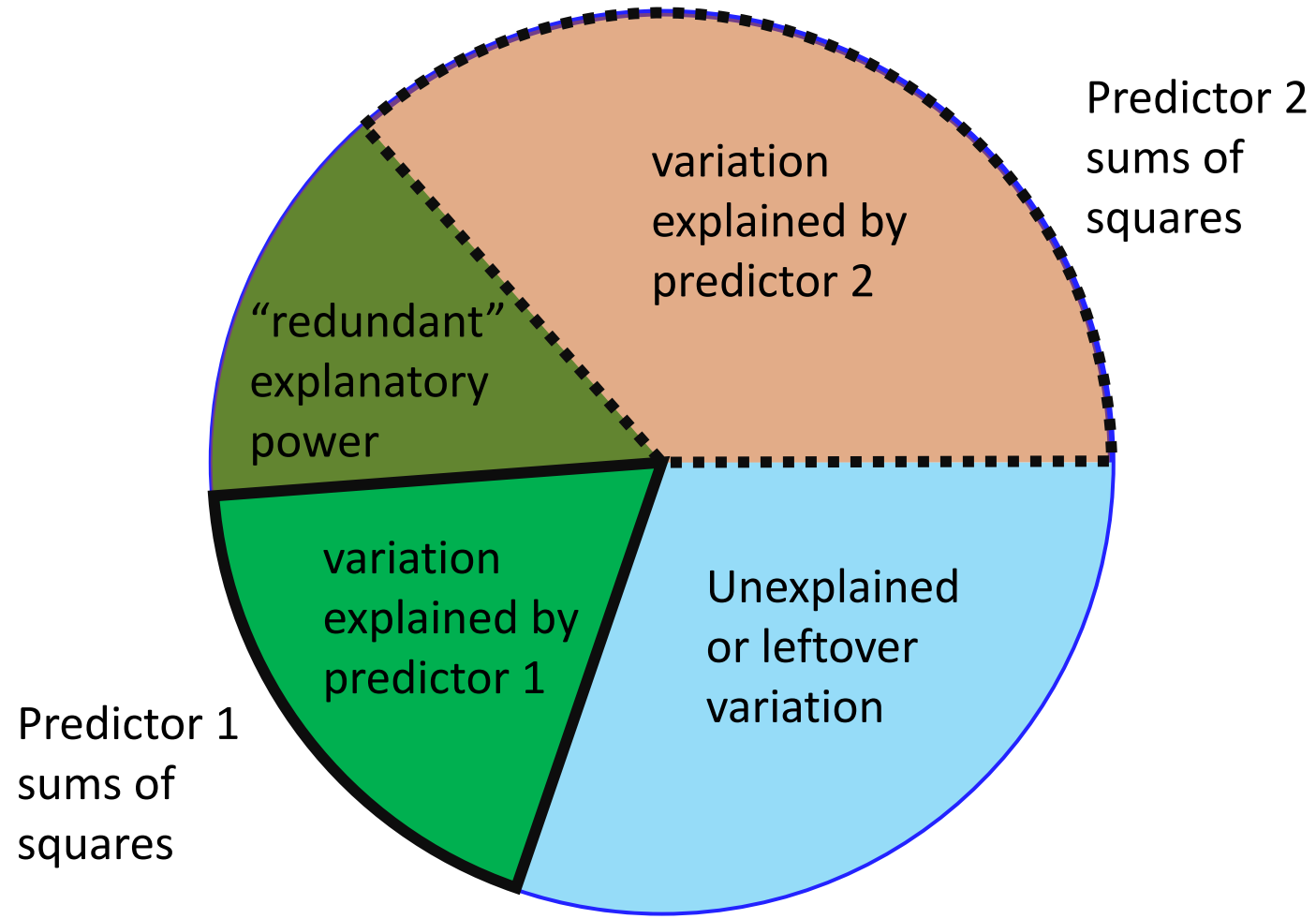
```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

# Sequential fit



Sequential fits change their answer depending on which variable/predictor is fit first.

# Marginal fit



# Sequential vs. marginal fitting

Table 1: ANOVA Table for fit\_ex1\_df (sequential fit)

Predictor	DF	SS	MS	F	P
f_YEAR	2	7049.72	3524.86	24.99	<0.0001
HEIGHT f_YEAR	1	6091.98	6091.98	43.2	<0.01
Residuals	399	56268.21	141.02		

Table 2: ANOVA Table for fit\_ex2\_df (sequential fit)

Predictor	DF	SS	MS	F	P
HEIGHT	1	7692.19	7692.19	54.55	<0.0001
f_YEAR HEIGHT	2	5449.52	2724.76	19.32	<0.0001
Residuals	399	56268.21	141.02		

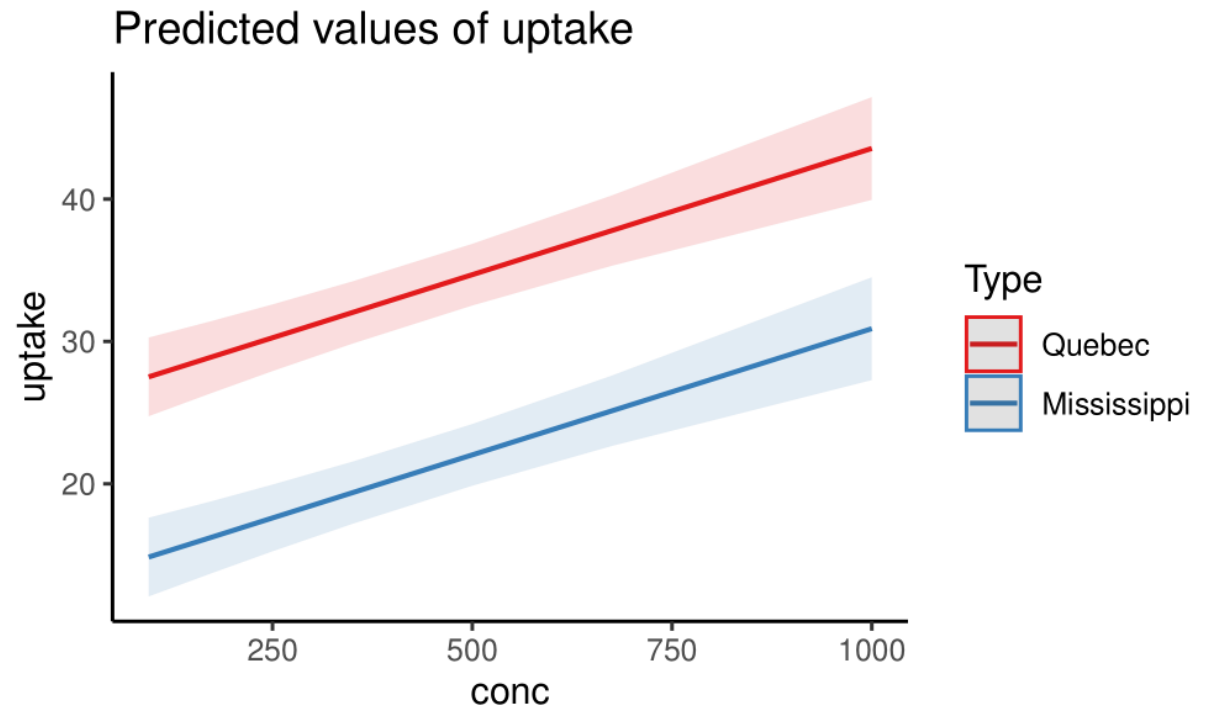
Table 3: ANOVA Table for fit\_ex1\_df (marginal fit)

Predictor	DF	SS	MS	F	P
f_YEAR HEIGHT	2	5449.52	2724.7600	19.32	<0.0001
HEIGHT f_YEAR	1	6091.98	6091.9800	43.2	<0.01
Residual	399	56268.21	141.0231		

# Main effects

```
fit_plants_1_nointeraction <- lm(uptake~conc+Type,data=C02)
summary(fit_plants_1_nointeraction)
```

```
##
## Call:
## lm(formula = uptake ~ conc + Type, data = C02)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -18.2145  -4.2549   0.5479   5.3048  12.9968
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    25.83005     1.579918  16.349  < 2e-16 ***
## conc           0.017731     0.002625   6.755 2.00e-09 ***
## TypeMississippi -12.659524     1.544261  -8.198 3.06e-12 ***
```



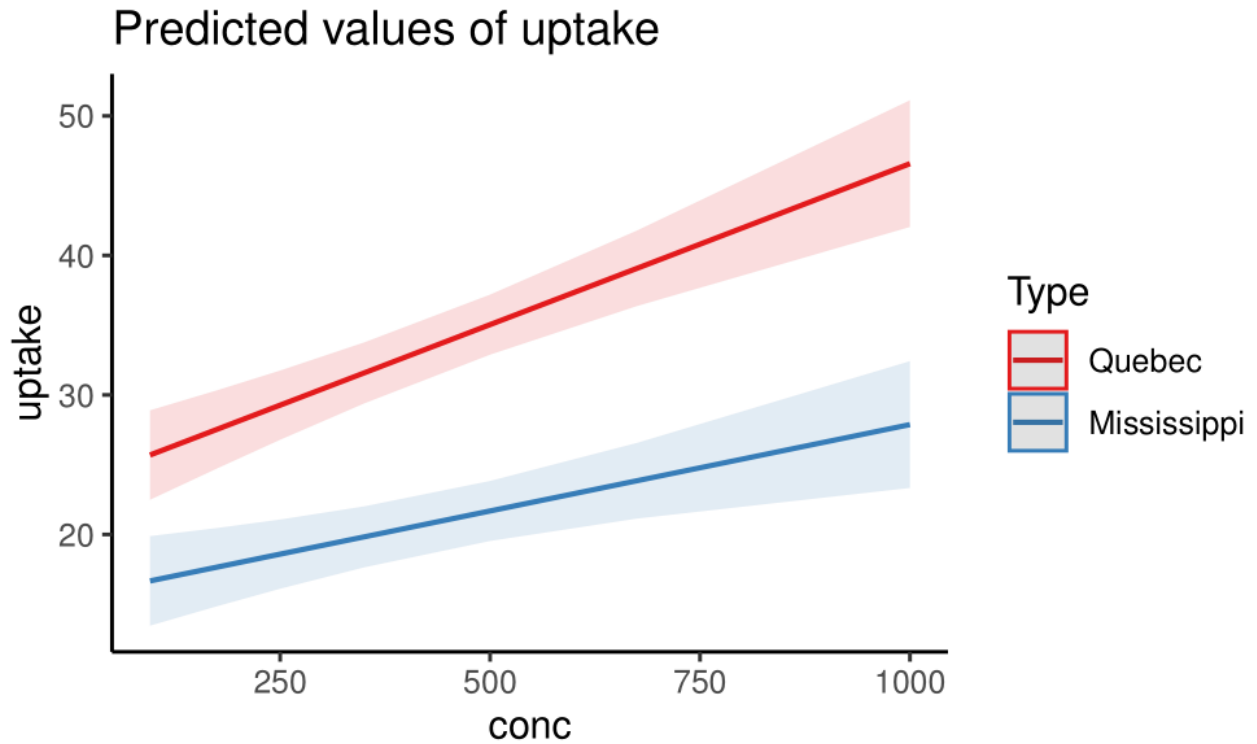
# Interactions

```
fit_plants_1_interaction <- lm(uptake~conc*Type,data=C02)
```

```
summary(fit_plants_1_interaction)
```

```
##
## Call:
## lm(formula = uptake ~ conc * Type, data = C02)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -16.3956  -5.5250  -0.1604   5.5724  12.0072
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    23.50303      1.91053   12.302 < 2e-16 ***
## conc           0.023080     0.003638    6.344 1.25e-08 ***
## TypeMississippi -8.005495     2.701899   -2.963 0.00401 **
## conc:TypeMississippi -0.010699     0.005145   -2.079 0.04079 *
##
```

$\text{lm}(\text{uptake} \sim \text{conc} + \text{Type} + \text{conc} * \text{Type}, \text{data} = \text{C02})$



# Interactions

```
fit_plants_1_interaction <- lm(uptake~conc*Type,data=C02)
```

```
summary(fit_plants_1_interaction)
```

```
##
## Call:
## lm(formula = uptake ~ conc * Type, data = C02)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -16.3956  -5.5250  -0.1604   5.5724  12.0072
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    23.50303      1.91053   12.302 < 2e-16 ***
## conc           0.023080     0.003638    6.344 1.25e-08 ***
## TypeMississippi -8.005495     2.701899   -2.963 0.00401 **
## conc:TypeMississippi -0.010699     0.005145   -2.079 0.04079 *
##
```

```
> Anova(fit_plants_1_interaction, type = "III")
```

Anova Table (Type III tests)

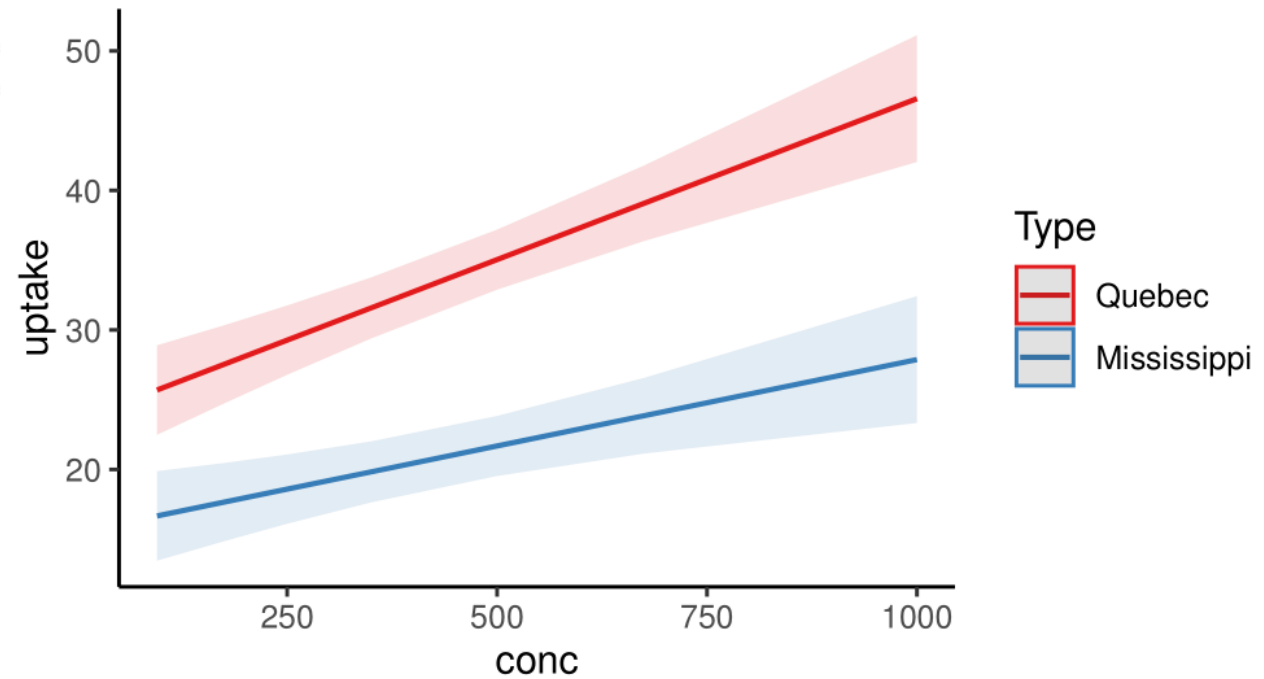
Response: uptake

	Sum Sq	Df	F value	Pr(>F)
(Intercept)	7280.1	1	151.3352	< 2.2e-16 ***
conc	1935.9	1	40.2426	1.253e-08 ***
Type	422.3	1	8.7789	0.004012 **
conc:Type	208.0	1	4.3238	0.040787 *
Residuals	3848.4	80		

---  
signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

$\text{lm}(\text{uptake} \sim \text{conc} + \text{Type} + \text{conc} * \text{Type}, \text{data} = \text{C02})$

Predicted values of uptake





# Evolutionary and plastic variation in larval growth and digestion reveal the complex underpinnings of size and age at maturation in dung beetles

Patrick T. Rohner  | Armin P. Moczek 


## Example multiple linear regression



Explore data | Search

Who we are | What we do | Join us | Help | Login

### Evolutionary and plastic variation in larval growth and digestion reveal the complex underpinnings of size and age at maturation in dung beetles

Rohner, Patrick T., Indiana University Bloomington,  <https://orcid.org/0000-0002-9840-1050>

Moczek, Armin, Indiana University Bloomington

[patrick.t.rohner@gmail.com](mailto:patrick.t.rohner@gmail.com)

Published Feb 16, 2023 on Dryad. <https://doi.org/10.5061/dryad.j9kd51cdc>


#### Cite this dataset

Rohner, Patrick T.; Moczek, Armin (2023). Evolutionary and plastic variation in larval growth and digestion reveal the complex underpinnings of size and age at maturation in dung beetles [Dataset]. Dryad. <https://doi.org/10.5061/dryad.j9kd51cdc>

#### Abstract

Age and size at maturity are key life history components, yet the proximate underpinnings that mediate intra- and interspecific variation in life history remain poorly understood. We studied the proximate underpinnings of species differences and nutritionally plastic variation in adult size and development time in four species of

#### Data files


 Download dataset


> Oct 21, 2022  
> Jan 21, 2023


#### Share



#### Metrics

 28 views

 3 downloads

 0 citations

### *Liatongus militaris*

