## Two Lines: Crocodiles!!

## head(crocs)

| \#\# | Body | Head | Species |
| ---: | ---: | ---: | ---: |
| \#\# | 1 | 338.0 | 52.0 |$\quad$ Indian

ggplot (data $=$ crocs, mapping $=$ aes $(x=$ Body, $y=$ Head, color $=$ Species $)$ ) + geom_point() + geom_smooth(method = "lm", se = FALSE)


2 lines by filtering to create separate data sets

```
aus_crocs <- crocs %>% filter(Species == "Australian")
aus_fit <- lm(Head ~ Body, data = aus_crocs)
summary(aus_fit)
##
## Call:
## lm(formula = Head ~ Body, data = aus_crocs)
##
## Residuals:
\begin{tabular}{lrrrrr} 
\#\# & Min & 1Q & Median & 3Q & Max \\
\#\# & -2.3529 & -0.9968 & 0.0824 & 0.7419 & 2.7973
\end{tabular}
##
## Coefficients:
```

```
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.463022 1.523732 2.273 0.0407 *
## Body 0.125344 0.004819 26.010 1.35e-12 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.504 on 13 degrees of freedom
## Multiple R-squared: 0.9811, Adjusted R-squared: 0.9797
## F-statistic: 676.5 on 1 and 13 DF, p-value: 1.35e-12
ind_crocs <- crocs %>% filter(Species == "Indian")
ind_fit <- lm(Head ~ Body, data = ind_crocs)
summary(ind_fit)
```

\#\#
\#\# Call:
\#\# lm(formula = Head ~ Body, data = ind_crocs)
\#\#
\#\# Residuals:

| \#\# | Min | 1Q | Median | 3Q | Max |
| :--- | ---: | ---: | ---: | ---: | ---: |
| \#\# | -4.5756 | -1.6627 | -0.0904 | 1.2208 | 4.6261 |

\#\#
\#\# Coefficients:
\#\# Estimate Std. Error t value $\operatorname{Pr}(>|t|)$
\#\# (Intercept) 10.538438 1.861787 5.66 4.53e-05 ***
\#\# Body $0.1313040 .005791 \quad 22.685 .08 \mathrm{e}-13$ ***
\#\# ---
\#\# Signif. codes: $0{ }^{\prime} * * * ' 0.001{ }^{\prime} * * ' 0.01{ }^{\prime} *^{\prime} 0.05{ }^{\prime} .{ }^{\prime} 0.1$ ' ' 1
\#\#
\#\# Residual standard error: 2.503 on 15 degrees of freedom
\#\# Multiple R-squared: 0.9717, Adjusted R-squared: 0.9698
\#\# F-statistic: 514.2 on 1 and 15 DF, p-value: $5.08 \mathrm{e}-13$

Questions we'd like to be able to answer (but can't with this output):

1. Is there statistically significant evidence that the intercepts for these lines are different?
2. Is there statistically significant evidence that the slopes for these lines are different?

## 2 parallel lines (same slope)

- Our Goal: Equations for two lines

Predicted Head Length for Australian Crocs $=\hat{\beta}_{0}^{\text {Australian }}+\hat{\beta}_{1} \times$ (Body Length $)$ Predicted Head Length for Indian Crocs $=\hat{\beta}_{0}^{\text {Indian }}+\hat{\beta}_{1} \times($ Body Length $)$

- Note: Different intercepts, same slope.

```
parallel_lines_fit <- lm(Head ~ Body + Species, data = crocs)
summary(parallel_lines_fit)
```

```
##
## Call:
## lm(formula = Head ~ Body + Species, data = crocs)
##
## Residuals:
\begin{tabular}{lrrrrr} 
\#\# & Min & \(1 Q\) & Median & 3Q & Max \\
\#\# & -4.4959 & -1.4218 & -0.0842 & 1.0117 & 4.6405
\end{tabular}
##
## Coefficients:
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.265418 1.309167 1.73 0.0942 .
## Body 0.129261 0.003904 33.11 < 2e-16 ***
## SpeciesIndian 8.893772 0.737538 12.06 8.05e-13 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.082 on 29 degrees of freedom
## Multiple R-squared: 0.977, Adjusted R-squared: 0.9755
## F-statistic: }617\mathrm{ on 2 and 29 DF, p-value: < 2.2e-16
```

- R gives us a single combined equation:

Predicted Head Length $=\hat{\beta}_{0}+\hat{\beta}_{1}($ Body Length $)+\hat{\beta}_{2}$ SpeciesIndian

Predicted Head Length $=2.27+0.13($ Body Length $)+8.89$ SpeciesIndian

## What is the SpeciesIndian variable?

- Behind the scenes, R creates a new indicator variable called SpeciesIndian:

$$
\text { SpeciesIndian }= \begin{cases}1 & \text { if the species for crocodile } i \text { is Indian. } \\ 0 & \text { otherwise (in this case, the species is Australian) }\end{cases}
$$

- R doesn't modify the data frame (it creates a secret copy in the background), but it would look like this:

```
head(crocs)
## Body Head Species SpeciesIndian
## 1 338.0 52.0 Indian 1
## 2 333.0 48.0 Australian 0
## 3 202.0 38.3 Indian 1
## 4 406.0 52.0 Australian 0
## 5 459.4 60.5 Australian 0
## 6 264.0 49.0 Indian 1
```

Above, we obtained this estimated equation:

Predicted Head Length $=2.27+0.13($ Body Length $)+8.89$ SpeciesIndian

What is the estimated equation describing the relationship between body length and head length, for Australian crocodiles?

What is the estimated equation describing the relationship between body length and head length, for Indian crocodiles?

What is the interpretation of $\widehat{\beta}_{0}=2.27$ ?

What is the interpretation of $\widehat{\beta}_{1}=0.13 ?$

What is the interpretation of $\widehat{\beta}_{2}=8.89 ?$

Using the output from the summary function, conduct a test of the claim that the intercept of the line describing the relationship between body length and head length in the population of all Australian crocodiles is the same as the intercept of the line describing the relationship between body length and head length in the population of all Indian crocodiles.

## Lines with different slopes (interactions)

To get different slopes, use Body * Species instead of Body + Species: two_lines_fit <- lm(Head ~ Species * Body, data = crocs) summary (two_lines_fit)

```
##
## Call:
## lm(formula = Head ~ Species * Body, data = crocs)
##
## Residuals:
## Min 1Q Median 3Q Max
## -4.5756 -1.3294 -0.0040 0.9646 4.6261
##
## Coefficients:
\begin{tabular}{lrrrr} 
\#\# & Estimate & Std. Error & t value & \(\operatorname{Pr}(>|\mathrm{t}|)\) \\
\#\# (Intercept) & 3.463022 & 2.126572 & 1.628 & 0.1146 \\
\#\# SpeciesIndian & 7.075415 & 2.638253 & 2.682 & \(0.0121 *\) \\
\#\# Body & 0.125344 & 0.006726 & 18.637 & \(<2 \mathrm{e}-16 * * *\) \\
\#\# SpeciesIndian:Body & 0.005959 & 0.008296 & 0.718 & 0.4785
\end{tabular}
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.099 on 28 degrees of freedom
## Multiple R-squared: 0.9775, Adjusted R-squared: 0.975
## F-statistic: 404.6 on 3 and 28 DF, p-value: < 2.2e-16
```

What is the estimated equation from this model?

What is the estimated equation describing the relationship between body length and head length, for Australian crocodiles?

What is the estimated equation describing the relationship between body length and head length, for Indian crocodiles?

What is the interpretation of $\widehat{\beta}_{0}=3.463$ ?

What is the interpretation of $\widehat{\beta}_{1}=7.075$ ?

What is the interpretation of $\widehat{\beta}_{2}=0.125$ ?

What is the interpretation of $\widehat{\beta}_{3}=0.006 ?$

Using the output from the summary function, conduct a test of the claim that the slope of the line describing the relationship between body length and head length in the population of all Australian crocodiles is the same as the slope of the line describing the relationship between body length and head length in the population of all Indian crocodiles.

