Salamanders

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Import libraries

```
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr 1.1.2 v readr 2.1.4
## v forcats 1.0.0 v stringr 1.5.0
## v ggplot2 3.4.3 v tibble 3.2.1
## v lubridate 1.9.2 v tidyr 1.3.0
## v purrr 1.0.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library(tidymodels)
## -- Attaching packages ------ tidymodels 1.1.1 --
## v broom 1.0.5 v rsample 1.2.0
## v dials 1.2.0 v tune 1.1.2
## v infer 1.0.5 v workflows 1.1.3
## v modeldata 1.2.0 v workflowsets 1.0.1
## v parsnip 1.1.1 v yardstick 1.2.0 ## v recipes 1.0.8
## -- Conflicts ----- tidymodels_conflicts() --
## x scales::discard() masks purrr::discard()
## x dplyr::filter() masks stats::filter()
## x recipes::fixed() masks stringr::fixed()
## x dplyr::lag() masks stats::lag()
## x yardstick::spec() masks readr::spec()
## x recipes::step() masks stats::step()
## * Use suppressPackageStartupMessages() to eliminate package startup messages
library(ggforce)
library(yardstick)
```

Import Salamanders data

```
salamanders <- read_csv("Salamanders.csv") %>% as_tibble()

## Rows: 11 Columns: 2

## -- Column specification ------

## Delimiter: ","

## dbl (2): RockDensity, SalamanderDensity

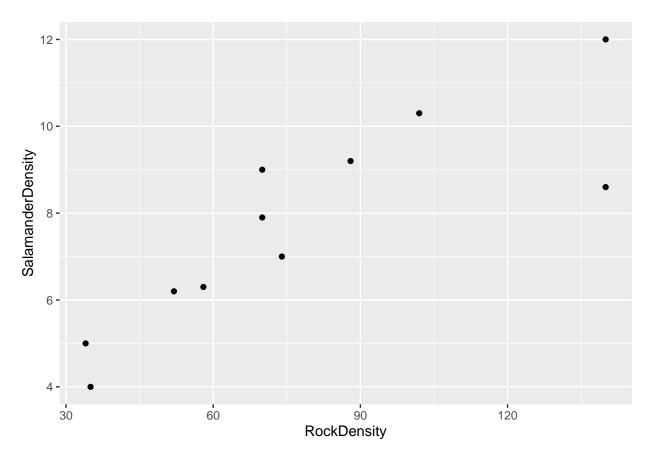
##

## i Use 'spec()' to retrieve the full column specification for this data.
```

Create a plot

```
ggplot(data = salamanders) + geom_point(aes(x = RockDensity, y = SalamanderDensity))
```

i Specify the column types or set 'show_col_types = FALSE' to quiet this message.



Create a linear regression model

```
model <- lm(SalamanderDensity ~ RockDensity, data = salamanders)
model

##
## Call:
## lm(formula = SalamanderDensity ~ RockDensity, data = salamanders)
##
## Coefficients:
## (Intercept) RockDensity
## 3.458 0.055</pre>
```

```
summary(model)
##
## lm(formula = SalamanderDensity ~ RockDensity, data = salamanders)
## Residuals:
      Min
               1Q Median 3Q
                                      Max
## -2.5576 -0.4378 -0.1178 0.8724 1.6923
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.45792 0.96599 3.580 0.005934 **
## RockDensity 0.05500
                          0.01125 4.888 0.000862 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 1.301 on 9 degrees of freedom
## Multiple R-squared: 0.7264, Adjusted R-squared: 0.696
## F-statistic: 23.89 on 1 and 9 DF, p-value: 0.0008621
salamanders$sqrtSalamanderDensity <- salamanders$SalamanderDensity^0.5
model2 <- lm(sqrtSalamanderDensity ~ RockDensity, data = salamanders)</pre>
model2
##
## lm(formula = sqrtSalamanderDensity ~ RockDensity, data = salamanders)
##
## Coefficients:
## (Intercept) RockDensity
      1.97228
                 0.01001
summary(model2)
##
## lm(formula = sqrtSalamanderDensity ~ RockDensity, data = salamanders)
## Residuals:
##
       Min
                 1Q Median
                                   3Q
## -0.44077 -0.07178 -0.00270 0.15904 0.32719
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.972285
                         0.179183 11.007 1.6e-06 ***
## RockDensity 0.010008 0.002087 4.795 0.000981 ***
## ---
```

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

Residual standard error: 0.2413 on 9 degrees of freedom

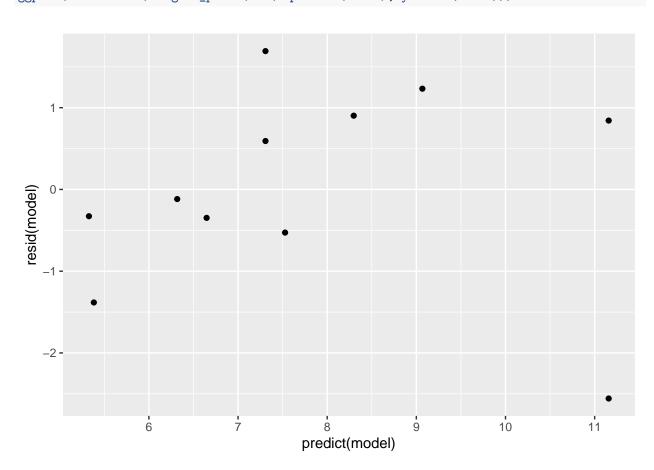
```
## Multiple R-squared: 0.7187, Adjusted R-squared: 0.6874
## F-statistic: 22.99 on 1 and 9 DF, p-value: 0.0009808
salamanders$SalamanderDensity2 <- salamanders$SalamanderDensity^2</pre>
model3 <- lm(SalamanderDensity2 ~ RockDensity, data = salamanders)</pre>
model3
##
## Call:
## lm(formula = SalamanderDensity2 ~ RockDensity, data = salamanders)
## Coefficients:
## (Intercept) RockDensity
##
      -2.9275
                    0.8719
summary(model3)
##
## Call:
## lm(formula = SalamanderDensity2 ~ RockDensity, data = salamanders)
## Residuals:
##
      Min
               1Q Median
                               ЗQ
                                      Max
## -45.176 -9.770 -1.716 15.464 24.864
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.9275 15.8443 -0.185 0.85751
## RockDensity 0.8719
                          0.1846 4.724 0.00108 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 21.34 on 9 degrees of freedom
## Multiple R-squared: 0.7126, Adjusted R-squared: 0.6807
## F-statistic: 22.32 on 1 and 9 DF, p-value: 0.001083
salamanders$reSalamanderDensity <- 1/salamanders$SalamanderDensity
model4 <- lm(reSalamanderDensity ~ RockDensity, data = salamanders)</pre>
model4
##
## Call:
## lm(formula = reSalamanderDensity ~ RockDensity, data = salamanders)
## Coefficients:
## (Intercept) RockDensity
      0.22542
                  -0.00107
```

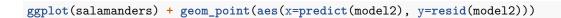
summary(model4)

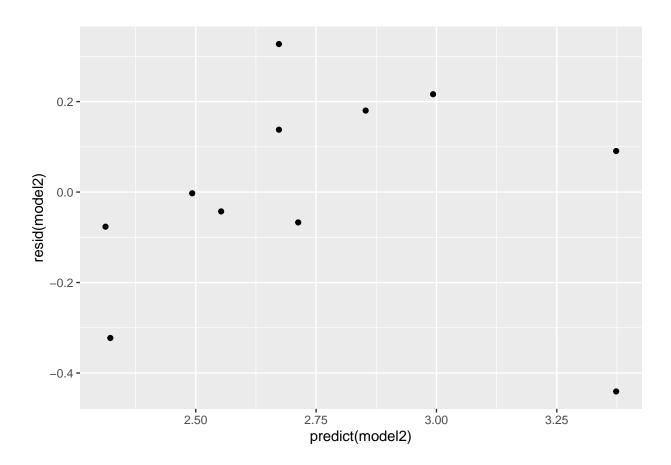
```
##
## Call:
## lm(formula = reSalamanderDensity ~ RockDensity, data = salamanders)
##
## Residuals:
##
        Min
                   1Q
                         Median
                                       3Q
                                                Max
  -0.039389 -0.020853 -0.004612 0.009361 0.062043
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 0.2254153 0.0232048 9.714 4.55e-06 ***
## RockDensity -0.0010702 0.0002703 -3.959 0.00331 **
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 0.03125 on 9 degrees of freedom
## Multiple R-squared: 0.6353, Adjusted R-squared: 0.5948
## F-statistic: 15.68 on 1 and 9 DF, p-value: 0.003307
```

Model 1 is the most accurate, as it has the lowest r^2 value

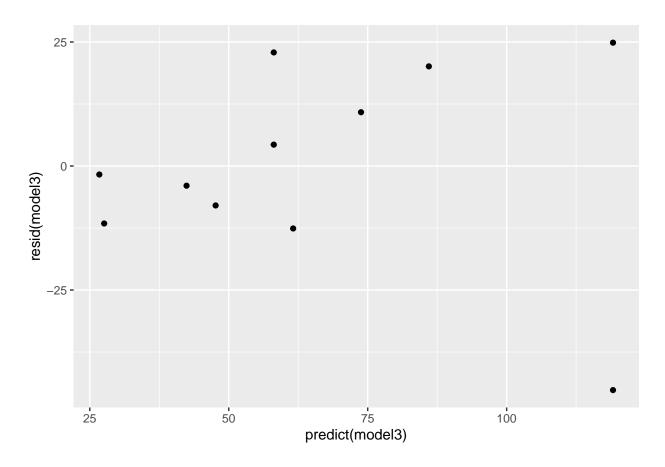
ggplot(salamanders) + geom_point(aes(x=predict(model), y=resid(model)))







ggplot(salamanders) + geom_point(aes(x=predict(model3), y=resid(model3)))



ggplot(salamanders) + geom_point(aes(x=predict(model4), y=resid(model4)))

