



Multiple Linear Regression in R

Fitting Models to Data Not Data to Models

Model Fitting Series - With Applications in R

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October 16, 2025

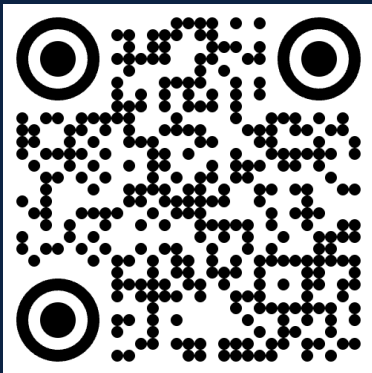
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Install Required Packages

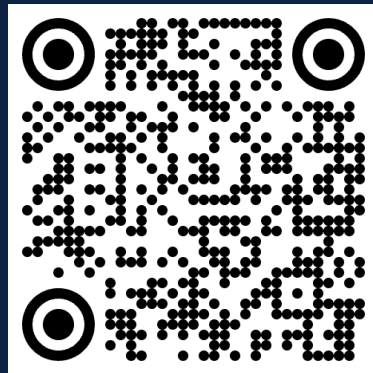
```
packages <- c("dplyr", "mgcv", "ggplot2",  
             "gratia", "dagitty", "leaps", "car")  
toInstall <- packages[!(packages %in%  
                        installed.packages()[,"Package"])]  
if(length(toInstall)) install.packages(toInstall)  
  
library('dplyr')    # for data wrangling  
library('mgcv')     # for modeling  
library('ggplot2')  # for fancy plots  
library('dagitty')  # for DAGs  
library('gratia')   # ggplot-based graphics  
library('leaps')    # for best subset regression  
library('car')      # for variance inflation factors
```

Session	Topic	Date/Time
1	Simple Linear Regression	Oct 7, 9:00 AM
2	Fitting Linear Models in R	Oct 8, 10:30 AM
3	Multiple Linear Regression in R	Oct 16, 4:00 PM
4	Interaction Terms & Hierarchical Linear Models	Oct 21, 11:00 AM
5	Generalized Linear Models	Oct 23, 4:00 PM
6	Generalized Additive Models (GAMs)	Oct 28, 11:00 AM
7	Interpreting & Predicting from GAMs	Oct 29, 10:30 AM
8	Hierarchical GAMs	Nov 4, 12:00 PM
9	Penalized Models	Nov 18, 11:00 AM
10	Survival Models	Nov 25, 11:00 AM
11	Nonparametric Models	Dec 2, 11:00 AM



←
New to R? Check
out the Fundamen-
tals of R series!

GitHub code and
slides for today's
workshop (and pre-
vious workshops)



Alternatively, code/slides available at the bottom of
<https://csc-ubc-okanagan.github.io/workshops/>



Key Concepts

- Built a diagnostics function to check all 5 assumptions
- Fitted models to real datasets: `ChickWeight`, `prostate`, `state.x77`
- Saw how outliers can dominate inference, skew predictions.

Today: Multiple Linear Regression in R



Today we will...

- Extend from simple ($Y = \beta_0 + \beta_1 x + \epsilon$) to multiple regression
- Use DAGs (Directed Acyclic Graphs) to visualize relationships
- Interpret coefficients when predictors are correlated
- Understand R^2_{adj} and model selection
- Work with `state.x77` data: Life expectancy modeling

Today we require...

```
library('dplyr')    # for data wrangling
library('mgcv')     # for modeling
library('ggplot2')  # for fancy plots
library('gratia')   # for ggplot-based model graphics
library('dagitty')  # for drawing Directed Acyclical Graphs (DAGs)
library('leaps')    # for best subset regression
library('car')      # for variation inflation factors
```

From Simple to Multiple

Simple Linear Regression:

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

Multiple Linear Regression:

$$Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \cdots + \beta_p x_{ip} + \epsilon_i$$

where:

- Y_i is the response for observation i
- x_{ij} is the value of predictor j for observation i
- β_j is the coefficient for predictor j
- $\epsilon_i \sim N(0, \sigma^2)$ (same assumptions as before)

Least Squares Estimates: SLR vs MLR



Simple Linear Regression

Model: $Y = \beta_0 + \beta_1 x + \epsilon$ Estimates:

$$\hat{\beta}_1 = \frac{\sum_{i=1}^n (x_i - \bar{x})(Y_i - \bar{Y})}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

$$\hat{\beta}_0 = \bar{Y} - \hat{\beta}_1 \bar{x}$$

Multiple Linear Regression

Model: $Y = \beta_0 + \beta_1 x_1 + \dots + \beta_p x_p + \epsilon$

Matrix form: $\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \epsilon$ Estimates:

$$\hat{\boldsymbol{\beta}} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{Y},$$

where \mathbf{X} is the model matrix.

Estimates for both are solved via least squares, but MLR requires matrix derivatives. In MLR, $\hat{\beta}_j$ is the effect of x_j holding all other predictors constant, while in SLR, $\hat{\beta}_1$ is the total effect of x on Y

Preparing state.x77 data

```
states <- state.x77 %>%  
  as.data.frame() %>%  
  mutate(State = rownames(.)) %>% # state.x77 has state names as row names  
  `rownames<-`(NULL) %>% # drop rownames  
  relocate(State, .before = 1) %>% # make the States column the first one  
  rename(Life_exp = `Life Exp`,  
         Murder_1e5 = Murder,  
         HS_grad_perc = `HS Grad`) %>%  
  as_tibble()
```

Variables include Life expectancy, Income (per capita, 1974), Murder rate (per 100k, 1976), High school graduation %

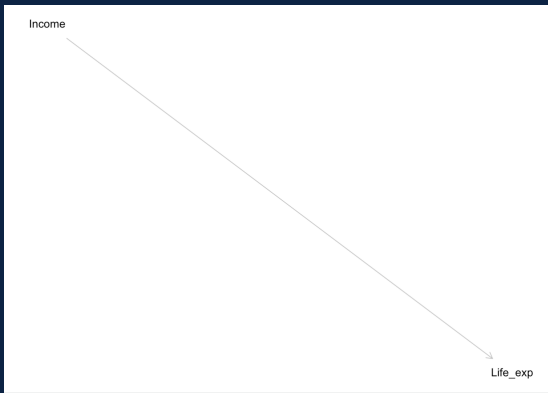
Directed Acyclic Graphs (DAGs)



What are DAGs?

- Visual representation of causal relationships
- Arrows show direction of causation
- Help identify confounders
- Guide model specification

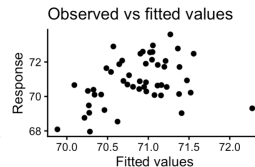
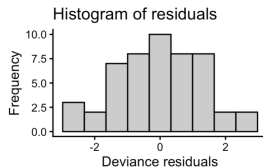
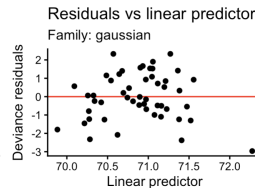
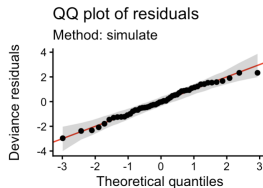
```
dag_1 <- dagitty('Income -> Life_exp')  
plot(dag_1)
```



Model 1: Income \rightarrow Life Expectancy



```
m_1 <- gam(Life_exp ~ Income,
            family = gaussian(),
            data = states,
            method = 'ML')
appraise(m_1, method = 'sim',
         n_simulate = 1e3)
```



Model 1: Income → Life Expectancy



```
summary(m_1) # relatively low  $R^2_{adj}$ 
```

Acceptable fit but relatively low R^2_{adj}
— suggests we might be missing important predictors

Formula:

Life_exp ~ Income

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	6.758e+01	1.328e+00	50.906	<2e-16 ***
Income	7.433e-04	2.965e-04	2.507	0.0156 *

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

R-sq.(adj) = 0.0974 Deviance explained = 11.6%
-ML = 82.089 Scale est. = 1.6266 n = 50

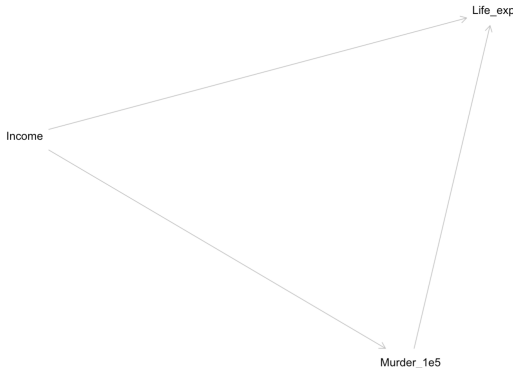
Model 2: Adding Murder Rate



DAG with two predictors

```
dag_2 <- dagitty(  
  'Income -> Life_exp  
    Murder_1e5 -> Life_exp  
    Income -> Murder_1e5')  
plot(dag_2)
```

Income affects life expectancy both directly AND indirectly through murder rate



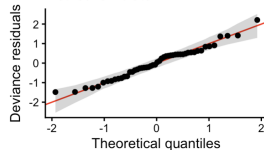
Model 2: Results I



```
m_2 <- gam(Life_exp ~ Income + Murder_1e5,  
            family = gaussian(),  
            data = states,  
            method = 'ML')  
appraise(m_2, method = 'sim',  
          n_simulate = 1e3)
```

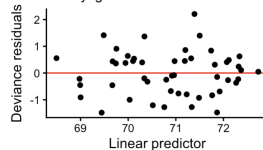
QQ plot of residuals

Method: simulate

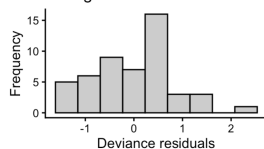


Residuals vs linear predictor

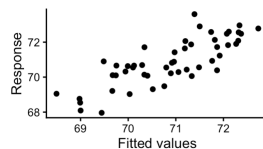
Family: gaussian



Histogram of residuals



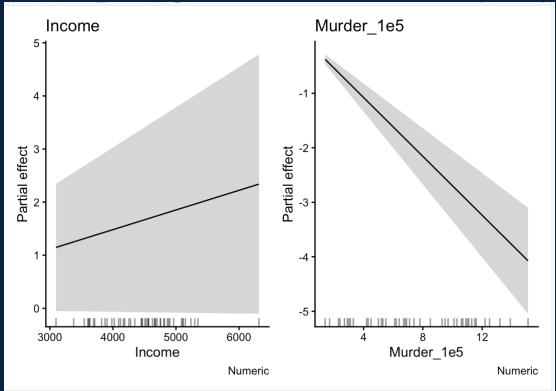
Observed vs fitted values



Model 2: Results II



```
draw(m_2, parametric = TRUE)
```



Model 2: Results III



```
summary(m_2) # relatively good R^2_adj  
# the p-value for Income went up. Why?
```

Formula:

Life_exp ~ Income + Murder_1e5

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	71.2255815	0.9673952	73.626	< 2e-16 ***
Income	0.0003705	0.0001973	1.878	0.0666 .
Murder_1e5	-0.2697594	0.0328408	-8.214	1.22e-10 ***

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

R-sq.(adj) = 0.622 Deviance explained = 63.7%

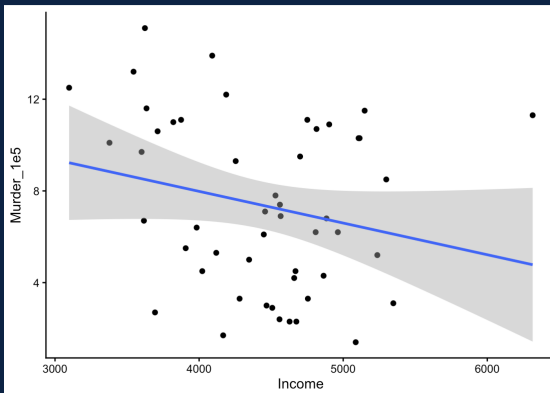
-ML = 59.834 Scale est. = 0.68205 n = 50

Understanding Multicollinearity



```
# murder rate and income are correlated,  
# but that's ok (see the DAG)  
ggplot(states, aes(Income, Murder_1e5)) +  
  geom_point() +  
  geom_smooth(method = 'lm',  
              formula = y ~ x)
```

When predictors are correlated,
coefficient interpretation changes:
each β represents the effect **holding
other variables constant**



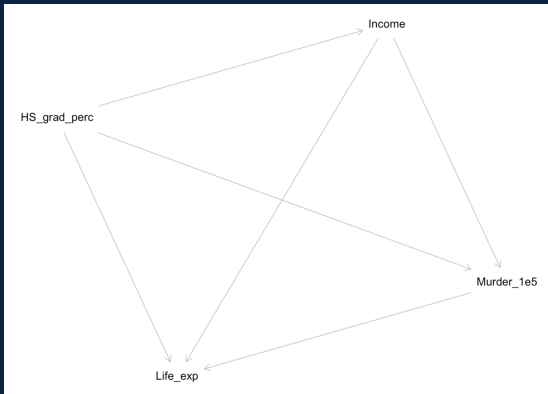
Model 3: Adding High School Graduation Rate



Complex DAG

```
dag_3 <- dagitty(  
  'Income -> Life_exp  
    Income -> Murder_1e5  
    Murder_1e5 -> Life_exp  
    HS_grad_perc -> Life_exp  
    HS_grad_perc -> Murder_1e5  
    HS_grad_perc -> Income')  
plot(dag_3)
```

Notice that HS graduation affects all other variables — a potential confounder!



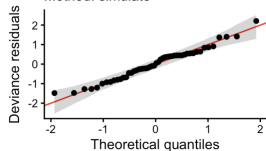
Model 3: Results I



```
m_3 <- gam(Life_exp ~ Income + Murder_1e5
            + HS_grad_perc,
            family = gaussian(),
            data = states,
            method = 'ML')
appraise(m_3, method = 'sim',
n_simulate = 1e3)
```

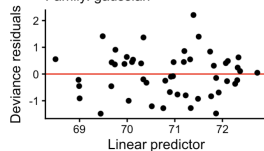
QQ plot of residuals

Method: simulate

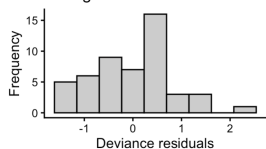


Residuals vs linear predictor

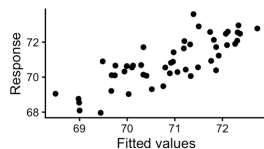
Family: gaussian



Histogram of residuals



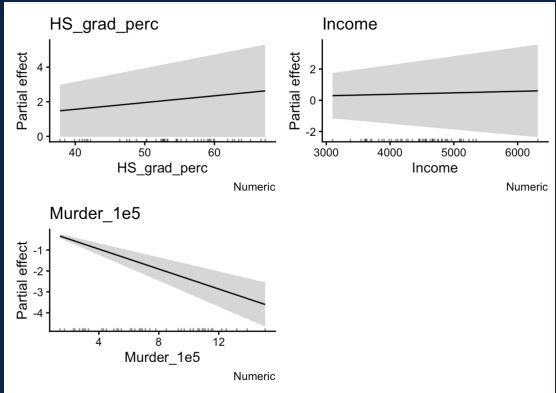
Observed vs fitted values



Model 3: Results II



```
draw(m_3, parametric = TRUE)
```



Model 3: Results III



```
summary(m_3) # R^2_adj did not improve much
```

Adding more predictors doesn't always improve the model!

Formula:

Life_exp ~ Income + Murder_1e5 + HS_grad_perc

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	7.014e+01	1.096e+00	63.979	< 2e-16 ***
Income	9.526e-05	2.393e-04	0.398	0.6924
Murder_1e5	-2.386e-01	3.581e-02	-6.664	2.92e-08 ***
HS_grad_perc	3.906e-02	2.030e-02	1.924	0.0605 .

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

R-sq.(adj) = 0.642 Deviance explained = 66.4%

-ML = 57.898 Scale est. = 0.64496 n = 50

Quick Sidenote on Model Matrix \mathbf{X}

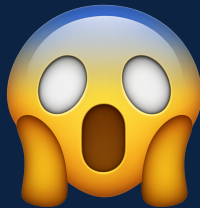


Recall that

$$\hat{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{Y},$$

where \mathbf{X} is an $n \times (p + 1)$ matrix, n is the number of observation in your data and p is the number of predictors (the $+1$ corresponds to an extra column for the intercept term β_0).

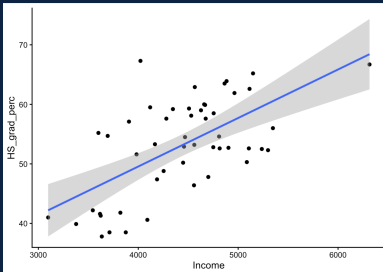
```
model_mat <- cbind(rep(1,nrow(states)),  
                  states$Income,  
                  states$Murder_1e5,  
                  states$HS_grad_perc)  
  
XTX_inv <- solve(t(model_mat)%*%model_mat)  
Betas <- XTX_inv%*%t(model_mat)%*%states$Life_exp  
Betas # Same estimates of previous slide!
```



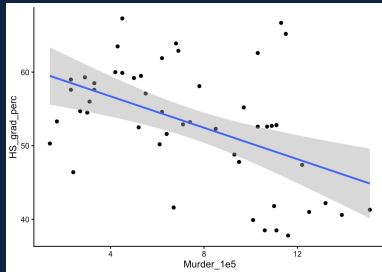
Correlation Structure



```
# % grad and income are correlated  
ggplot(states, aes(Income, HS_grad_perc)) +  
  geom_point() +  
  geom_smooth(method = 'lm', formula = y ~ x)
```



```
# same for % grad and murder rate  
ggplot(states, aes(Murder_1e5, HS_grad_perc)) +  
  geom_point() +  
  geom_smooth(method = 'lm', formula = y ~ x)
```



Caution High correlations between predictors make coefficients unstable and harder to interpret

VIFs

The VIF is given by

$$VIF_i = \frac{1}{1 - R_i^2},$$

which is used for each independent variable i in a multiple regression model. Here, R_i^2 is the R-squared value obtained from regressing that variable against all the other independent variables in the model.

```
m_3.1 <- lm(Life_exp~Income + Murder_1e5 + HS_grad_perc,
             data = states)
vif(m_3.1)
m_3.2 <- lm(Life_exp~.-State,
             data = states)
vif(m_3.2)
```

A $VIF \geq 10$ indicates severe multicollinearity, consider removing that variable, whereas $5 \leq VIF < 10$ should be investigated.

```
> vif(m_3.1)
      Income      Murder_1e5      HS_grad_perc
      1.642581      1.327381      2.041819
> vif(m_3.2)
      Population      Income      Illiteracy      Murder_1e5      HS_grad_perc
      1.499915      1.992680      4.403151      2.616472      3.134887
      Frost      Area
      2.358206      1.789764
>
```


Key Concepts

- **Partial effects:** Each β_j represents the effect of x_j *holding all other predictors constant*
- R_{adj}^2 : Adjusts for number of predictors: $R_{adj}^2 = 1 - \frac{(1-R^2)(n-1)}{n-p-1}$
- **Multicollinearity:** When predictors are correlated:
 - Standard errors increase
 - Coefficients become unstable
 - Individual p-values may be misleading

Statistical significance \neq Causal effect; use DAGs to think about causality.

Good Reasons to Add Variables

- Theory/domain knowledge suggests causality
- Substantial improvement in R^2_{adj}
- Reduces omitted variable bias
- Improves predictions on new data

Bad Reasons to Add Variables

- Just to increase R^2 (not adjusted)
- "Fishing" for significance
- Without theoretical justification
- When it creates severe multicollinearity

Next time: We'll explore interaction terms — when the effect of one variable depends on another.

A Handy Little Function



```
# part of the 'leaps' package
m_full <- regsubsets(Life_exp~State,
                    data = states,
                    nvmax = 5,
                    really.big = T)

summary(m_full)
best_ss <- gam(Life_exp~Murder_1e5,
              data = states,
              method = "ML")
summary(best_ss)
```

Well, turns out possibly the most obvious model is the "best" according to best subset regression

```
1 subsets of each size up to 5
Selection Algorithm: exhaustive
      Population Income Illiteracy Murder_1e5 HS_grad_perc Frost Area
1 ( 1 ) " " " " " " "*" " " " " " "
2 ( 1 ) " " " " " " "*" " " " " "
3 ( 1 ) " " " " " " "*" " " " " "
4 ( 1 ) "*" " " " " "*" " " " " "
5 ( 1 ) "*" "*" " " "*" " " " " "

> best_ss <- gam(Life_exp~Murder_1e5, data = states, method = "ML")
> summary(best_ss)

Family: gaussian
Link function: identity

Formula:
Life_exp ~ Murder_1e5

Parametric coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)  72.97356    0.26997   270.30 < 2e-16 ***
Murder_1e5   -0.28395    0.03279   -8.66 2.26e-11 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R-sq.(adj) =  0.602   Deviance explained =  61%
-ML = 61.642   Scale est. = 0.71794   n = 50
```

- Multiple regression extends simple regression: $Y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \cdots + \epsilon$
- DAGs help visualize and reason about causal relationships
- Correlated predictors complicate interpretation but are often unavoidable
- R^2_{adj} penalizes model complexity — more predictors isn't always better
- Each coefficient represents a *partial* effect, holding others constant
- Think causally, not just statistically!

Exercise: Choose a dataset and fit a MLR model

```
# Option 1: mtcars - Motor Trend Car Road Tests
?mtcars # Predict mpg by finding a good combination of predictors

# Option 2: airquality - New York Air Quality
?airquality # Predict Ozone by finding a good combination of predictors

# Option 3: swiss - Swiss Fertility and Socioeconomic Data
?swiss # Predict Fertility by finding a good combination of predictors

# Option 4: attitude - Chatterjee-Price Attitude Data
?attitude # Predict rating by finding a good combination of predictors

# 1. Draw a DAG for your chosen variables
# 2. Fit the model using gam()
# 3. Check diagnostics with appraise()
# 4. Interpret coefficients and  $R^2_{adj}$ 
# 5. Try adding/removing predictors - does it improve the model?
```

Additional Questions?
Book an Appointment!



Next Workshop

Interaction Terms & Hierarchical
Linear Models

October 21, 11:00 AM

- When effects depend on context
- Random effects
- Mixed models