#### Multiple Linear Regression



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#### Outline

Multiple Linear Regression

Model Fit and Model Comparison

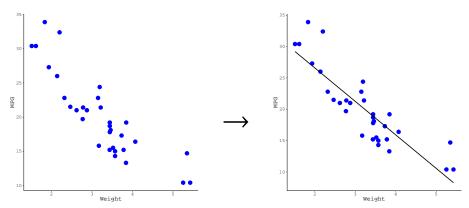
**Categorical Predictors** 



# Graphical Representations of Regression Models

A regression of two variables can be represented on a 2D scatterplot.

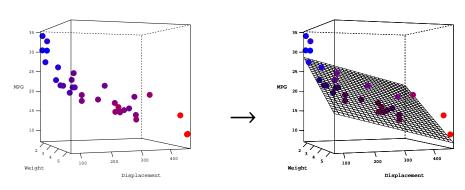
Simple linear regression implies a 1D line in 2D space.



# Graphical Representations of Regression Models

Adding an additional predictor leads to a 3D point cloud.

• A regression model with two IVs implies a 2D plane in 3D space.



#### **Partial Effects**

In MLR, we want to examine the *partial effects* of the predictors.

 What is the effect of a predictor after controlling for some other set of variables?

This approach is crucial to controlling confounds and adequately modeling real-world phenomena.



```
## Read in the 'diabetes' dataset:
dDat <- readRDS("../data/diabetes.rds")

## Simple regression with which we're familiar:
out1 <- lm(bp ~ age, data = dDat)</pre>
```

Asking: What is the effect of age on average blood pressure?



```
partSummary(out1, -1)
## Residuals:
##
      Min 1Q Median 3Q
                                   Max
## -31.188 -8.897 -1.209 8.612 39.952
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 77.47605 2.38132 32.535 < 2e-16
## age 0.35391 0.04739 7.469 4.39e-13
##
## Residual standard error: 13.04 on 440 degrees of freedom
## Multiple R-squared: 0.1125, Adjusted R-squared:
## F-statistic: 55.78 on 1 and 440 DF, p-value: 4.393e-13
```

```
## Add in another predictor:
out2 <- lm(bp ~ age + bmi, data = dDat)</pre>
```

Asking: What is the effect of BMI on average blood pressure, after controlling for age?

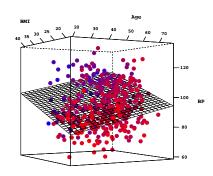
• We're partialing age out of the effect of BMI on blood pressure.



```
partSummary(out2, -1)
## Residuals:
      Min 1Q Median 3Q
                                   Max
##
## -29.287 -8.198 -0.178 8.413 41.026
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 52.24654 3.83168 13.635 < 2e-16
         0.28651 0.04504 6.362 5.02e-10
## age
## bmi
              1.08053 0.13363 8.086 6.06e-15
##
## Residual standard error: 12.18 on 439 degrees of freedom
## Multiple R-squared: 0.2276, Adjusted R-squared: 0.224
## F-statistic: 64.66 on 2 and 439 DF, p-value: < 2.2e-16
```

#### Interpretation

- The expected average blood pressure for an unborn patient with a negligible extent is 52.25.
- For each year older, average blood pressure is expected to increase by 0.29 points, after controlling for BMI.
- For each additional point of BMI, average blood pressure is expected to increase by 1.08 points, after controlling for age.



### Multiple $R^2$

How much variation in blood pressure is explained by the two models?

• Check the R<sup>2</sup> values.

```
## Extract R^2 values:
r2.1 <- summary(out1)$r.squared
r2.2 <- summary(out2)$r.squared
r2.1
## [1] 0.1125117
r2.2
## [1] 0.2275606</pre>
```

#### F-Statistic

How do we know if the  $R^2$  values are significantly greater than zero?

• We use the F-statistic to test  $H_0: R^2 = 0$  vs.  $H_1: R^2 > 0$ .

```
f1 <- summary(out1)$fstatistic
f1

## value numdf dendf
## 55.78116 1.00000 440.00000

pf(q = f1[1], df1 = f1[2], df2 = f1[3], lower.tail = FALSE)

## value
## 4.392569e-13</pre>
```

#### F-Statistic

```
f2 <- summary(out2)$fstatistic
f2

## value numdf dendf
## 64.6647 2.0000 439.0000

pf(f2[1], f2[2], f2[3], lower.tail = FALSE)

## value
## 2.433518e-25</pre>
```

#### **Comparing Models**

How do we quantify the additional variation explained by BMI, above and beyond age?

• Compute the  $\Delta R^2$ 

```
## Compute change in R^2:
r2.2 - r2.1
## [1] 0.115049
```

How do we know if  $\Delta R^2$  represents a significantly greater degree of explained variation?

• Use an F-test for  $H_0$ :  $\Delta R^2 = 0$  vs.  $H_1$ :  $\Delta R^2 > 0$ 

```
## Is that increase significantly greater than zero?
anova(out1, out2)

## Analysis of Variance Table

##
## Model 1: bp ~ age
## Model 2: bp ~ age + bmi

## Res.Df RSS Df Sum of Sq F Pr(>F)

## 1 440 74873

## 2 439 65167 1 9706.1 65.386 6.057e-15 ***

## ---

## Signif. codes:

## 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

#### **Model Comparison**

We can also compare models based on their prediction errors.

• For OLS regression, we usually compare MSE values.

```
mse1 <- MSE(y_pred = predict(out1), y_true = dDat$bp)
mse2 <- MSE(y_pred = predict(out2), y_true = dDat$bp)
mse1
## [1] 169.3963
mse2
## [1] 147.4367</pre>
```

In this case, the MSE for the model with *BMI* included is smaller.

• We should prefer the the larger model.

# **CATEGORICAL PREDICTORS**



#### **Categorical Predictors**

Most of the predictors we've considered thus far have been *quantitative*.

- Continuous variables that can take any real value in their range
- Interval or Ratio scaling

We often want to include grouping factors as predictors.

- These variables are qualitative.
  - Their values are simply labels.
  - There is no ordering of the categories.
  - Nominal scaling



#### How to Model Categorical Predictors

We need to be careful when we include categorical predictors into a regression model.

The variables need to be coded before entering the model

Consider the following indicator of major:

$$X_{maj} = \{1 = Law, 2 = Economics, 3 = Data Science\}$$

 What would happen if we naïvely used this variable to predict program satisfaction?

#### How to Model Categorical Predictors

```
mDat <- readRDS("../data/major_data.rds")
mDat[seq(25, 150, 25), ]

## sat majF majN
## 25  1.9  law  1
## 50  1.4  law  1
## 75  4.3 econ  2
## 100  4.1 econ  2
## 125  5.7  ds  3
## 150  5.1  ds  3

out1 <- lm(sat ~ majN, data = mDat)</pre>
```

# How to Model Categorical Predictors

```
partSummary(out1, -1)
## Residuals:
##
     Min 10 Median 30 Max
## -1.303 -0.313 -0.113 0.342 1.342
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) -0.33200 0.12060 -2.753 0.00664
## majN
        2.04500 0.05582 36.632 < 2e-16
##
## Residual standard error: 0.5582 on 148 degrees of freedom
## Multiple R-squared: 0.9007, Adjusted R-squared: 0.9
## F-statistic: 1342 on 1 and 148 DF, p-value: < 2.2e-16
```

### **Dummy Coding**

The most common way to code categorical predictors is *dummy coding*.

- A G-level factor must be converted into a set of G-1 dummy codes.
- Each code is a variable on the dataset that equals 1 for observations corresponding to the code's group and equals 0, otherwise.
- The group without a code is called the reference group.



#### Example Dummy Code

Let's look at the simple example of coding biological sex:

	sex	male	
1	female	0	
2	male	1	
3	male	1	
4	female	0	
5	male	1	
6	female	0	
7	female	0	
8	male	1	
9	female	0	
10	female	0	



### **Example Dummy Codes**

Now, a slightly more complex example:

	drink	juice	tea
1	juice	1	0
2	coffee	0	0
3	tea	0	1
4	tea	0	1
5	tea	0	1
6	tea	0	1
7	juice	1	0
8	tea	0	1
9	coffee	0	0
10	juice	1	0



#### **Using Dummy Codes**

To use the dummy codes, we simply include the G-1 codes as G-1 predictor variables in our regression model.

$$Y = \beta_0 + \beta_1 X_{male} + \varepsilon$$
  

$$Y = \beta_0 + \beta_1 X_{juice} + \beta_2 X_{tea} + \varepsilon$$

- The intercept corresponds to the mean of Y for the reference group.
- Each slope represents the difference between the mean of Y in the coded group and the mean of Y in the reference group.

First, an example with a single, binary dummy code:

```
## Read in some data:
cDat <- readRDS("../data/cars_data.rds")

## Fit and summarize the model:
out2 <- lm(price ~ mtOpt, data = cDat)</pre>
```

```
partSummary(out2, -1)
## Residuals:
## Min 10 Median 30
                                  Max
## -10.341 -6.338 -3.141 2.662 38.059
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 23.841 1.623 14.691 <2e-16
## mtOpt
        -6.603 2.004 -3.295 0.0014
##
## Residual standard error: 9.18 on 91 degrees of freedom
## Multiple R-squared: 0.1066, Adjusted R-squared:
## F-statistic: 10.86 on 1 and 91 DF, p-value: 0.001403
```

#### Interpretations

- The average price of a car without the option for a manual transmission is  $\hat{\beta}_0 = 23.84$  thousand dollars.
- The average difference in price between cars that have manual transmissions as an option and those that do not is  $\hat{\beta}_1 = -6.6$  thousand dollars.



Fit a more complex model:

```
out3 <- lm(price ~ front + rear, data = cDat)
partSummary(out3, -1)
## Residuals:
##
      Min 1Q Median 3Q
                                    Max
## -14.050 -6.250 -1.236 3.264 32.950
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 17.63000 2.76119 6.385 7.33e-09
## front
           -0.09418 2.96008 -0.032 0.97469
## rear 11.32000 3.51984 3.216 0.00181
##
## Residual standard error: 8.732 on 90 degrees of freedom
## Multiple R-squared: 0.2006, Adjusted R-squared: 0.1829
## F-statistic: 11.29 on 2 and 90 DF, p-value: 4.202e-05
```

#### Interpretations

- The average price of a four-wheel-drive car is  $\hat{\beta}_0 = 17.63$  thousand dollars.
- The average difference in price between front-wheel-drive cars and four-wheel-drive cars is  $\hat{\beta}_1 = -0.09$  thousand dollars.
- The average difference in price between rear-wheel-drive cars and four-wheel-drive cars is  $\hat{\beta}_2 = 11.32$  thousand dollars.



#### Include two sets of dummy codes:

#### Interpretations

- The average price of a four-wheel-drive car that does not have a manual transmission option is  $\hat{\beta}_0 = 21.72$  thousand dollars.
- After controlling for drive type, the average difference in price between cars that have manual transmissions as an option and those that do not is  $\hat{\beta}_1 = -5.84$  thousand dollars.
- After controlling for transmission options, the average difference in price between front-wheel-drive cars and four-wheel-drive cars is  $\hat{\beta}_2 = -0.26$  thousand dollars.
- After controlling for transmission options, the average difference in price between rear-wheel-drive cars and four-wheel-drive cars is  $\hat{\beta}_3 = 10.52$  thousand dollars.

For variables with only two levels, we can test the overall factor's significance by evaluating the significance of a single dummy code.

For variables with more than two levels, we need to simultaneously evaluate the significance of each of the variable's dummy codes.

```
summary(out4)$r.squared - summary(out2)$r.squared
## [1] 0.1767569
anova(out2, out4)
## Analysis of Variance Table
##
## Model 1: price ~ mtOpt
## Model 2: price ~ mtOpt + front + rear
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 91 7668.9
## 2 89 6151.6 2 1517.3 10.976 5.488e-05 ***
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

What about models where a single nominal factor is the only predictor?

We can compare back to an "intercept-only" model.

```
r2Diff <- summary(out3)$r.squared - summary(out0)$r.squared
r2Diff
## [1] 0.2006386
anova(out0, out3)
## Analysis of Variance Table
##
## Model 1: price ~ 1
## Model 2: price ~ front + rear
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 92 8584.0
## 2 90 6861.7 2 1722.3 11.295 4.202e-05 ***
## Signif. codes:
## 0 '***! 0.001 '**! 0.01 '*! 0.05 '.! 0.1 ' ! 1
```

We don't actually need to do the explicit model comparison, though.

```
r2Diff
## [1] 0.2006386
summary(out3)$r.squared
## [1] 0.2006386
anova(out0, out3)[2, "F"]
## [1] 11.29494
summary(out3)$fstatistic[1]
      value
## 11,29494
```

#### Conclusion

- Each variable in a regression model corresponds to a dimension in the data-space.
  - A regression model with P predictors implies a P-dimensional (hyper)-plane in (P + 1)-dimensional space.
- The coefficients in MLR are partial coefficients.
  - Each effect is interpreted as holding other predictors constant.
- Categorical predictors must be coded before they can be used in our models.
  - The regression coefficients represent group mean differences.