

MULTIPLE LINEAR REGRESSION



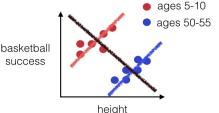
- Motivation
- Model and Interpretation
- Estimation and Inference
- Interaction



- The response or dependent variable, Y, may depend on several predictors not just one!
- Multiple regression is an attempt to consider the <u>simultaneous</u> influence of several variables on the response
- This may be with the goal of an unbiased estimate of *association* or for better *prediction*

Motivation

- Why not fit multiple separate simple linear regressions?
 - If the goal is to estimate the *association* between the response and a predictor of interest, a confounder can make the observed association appear
 - stronger than the true association,
 - weaker than the true association, or
 - even the reverse of the true association
- How can we address this:
 - We can adjust for the effects of the confounder by adding a corresponding term to our linear regression
- If the goal is *prediction* of the response, we may be able to improve prediction by including additional variables in the regression model



Motivation: Cholesterol Example

Data

>	> head(cholesterol)										
	ID	DM	age	chol	BMI	ΤG	APOE	rs174548	rs4775401	HTN	chd
	1	1	74	215	26.2	367	4	1	2	1	1
	2	1	51	204	24.7	150	4	2	1	1	1
	3	0	64	205	24.2	213	4	0	1	1	1
	4	0	34	182	23.8	111	2	1	1	1	0
	5	1	52	175	34.1	328	2	0	0	1	0
	6	1	39	176	22.7	53	4	0	2	0	0

- Our goal:
 - Investigate the relationship between age (years), BMI (kg/m²) and serum total cholesterol (mg/dl)

Motivation

In general, the multiple regression equation can be written as follows:

$$\mathsf{E}[\mathsf{Y} | \mathsf{x}_1, \mathsf{x}_2, \dots, \mathsf{x}_p] = \beta_0 + \beta_1 \mathsf{x}_1 + \beta_2 \mathsf{x}_2 + \dots + \beta_p \mathsf{x}_p$$

- We use multiple variables when:
 - The predictor variable is categorical with more than two groups
 - We need polynomials, splines or other functions to model the shape of the relationship(s) accurately
- Estimating association:
 - We want to adjust for confounding by other variables
 - We want to allow the association to differ for different values of other variables (interaction)
- Prediction: we use multiple variables if we think more than one variable will be useful in predicting future outcomes accurately

Model and Interpretation

• Model:
$$Y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + ... + \beta_p x_p + \varepsilon$$

where we assume $\varepsilon \sim N(0, \sigma^2)$

Extension of simple linear regression!

• Systematic component:

$$E[Y|x_1,...,x_p] = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + ... + \beta_p x_p$$

Random component:

$$Var[Y \mid x_1, ..., x_p] = \sigma^2$$

Model and Interpretation

• For example, let us assume that there are two predictors in the model and so $E[Y|x_1, x_2] = \beta_0 + \beta_1 x_1 + \beta_2 x_2$

Consider two observations with the same value for x_2 , but one observation has x_1 one unit higher, that is,

Obs 1:
$$E[Y|x_1=k+1, x_2=c] = \beta_0 + \beta_1 (k+1) + \beta_2 c$$

Obs 2: $E[Y|x_1=k, x_2=c] = \beta_0 + \beta_1 (k) + \beta_2 c$

Thus,
$$E[Y|x_1=k+1, x_2=c] - E[Y|x_1=k, x_2=c] = \beta_1$$

That is, β_1 is the expected mean change in y per unit change in x_1 if x_2 is held constant (adjusted/controlling for x_2)

Similar interpretation applies to β_2

Model and Interpretation

- To facilitate our discussion let's assume we have two predictors with binary values
- Model:

$$E[Y | x_1, x_2] = \beta_0 + \beta_1 x_1 + \beta_2 x_2$$

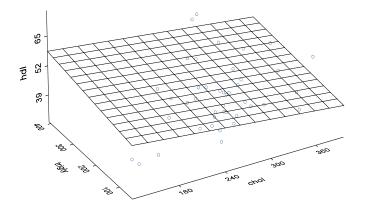
Mean of Y	X ₂ =0	X ₂ =1	$E[Y x_1=1, x_2=0] - E[Y x_1=0, x_2=0] = \beta_1$
X ₁ =0	βο	$\beta_0 + \beta_2$	$E[Y x_1=1, x_2=1] - E[Y x_1=0, x_2=1] = \beta_1$
$X_1 = 1$	$\beta_0 + \beta_1$	$\beta_0 + \beta_1 + \beta_2$	$E[Y x_1=0, x_2=1] - E[Y x_1=0, x_2=0] = \beta_2$
	P0 ' P1	P0 P1 P2	$E[Y x_1=1, x_2=1] - E[Y x_1=1, x_2=0] = \beta_2$

Estimation

- Least Squares Estimation:
 - As in linear regression, chooses the coefficient estimates that minimize the residual sum of squares

$$D = \sum_{i} (y_i - \hat{y}_i)^2$$

• Computation more difficult, but statistical software (R) will do that for you!



Estimation and Inference

- Inference
 - About regression model parameters
 - Hypothesis Testing H_0 : $\beta_j=0$ (j=0,1,2,...,p)

<u>Interpretation</u>: Is there a statistically significant relationship between the response y and x_j after adjusting for all other factors (predictors) in the model?

Test Statistic:

$$\frac{\hat{\beta}_j - (null \ hyp)}{se(\hat{\beta}_j)} \sim t_{n-p-1}$$

Note: The square of the t-statistic gives the F-statistic and the test is known as the **partial F-Test**

Confidence Intervals

$$\hat{\beta}_{j} \pm (critical \ value) \times se(\hat{\beta}_{j})$$

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Estimation and Inference

- About the full model
 - Hypotheses

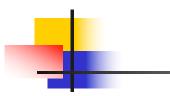
H₀: $\beta_1 = \beta_2 = ... = \beta_p = 0$ vs. H₁: At least one β_j is not null

Analysis of variance table

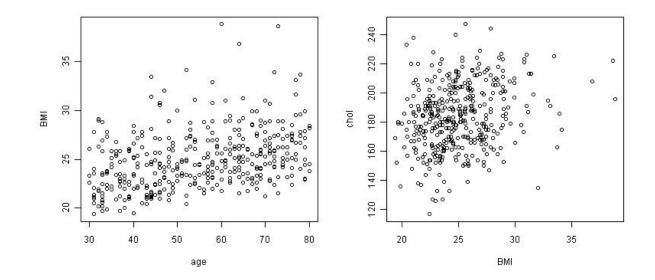
Source	df	SS	MS	F
Regression	р	$SSR = \sum (\hat{y}_i - \overline{y}_i)^2$	MSR= SSR/p	MSR/MSE
Residual	n-p-1	$SSE = \sum (y_i - \hat{y}_i)^2$	MSE= SSE/(n-p-1)	
Total	n-1	$SST = \sum (y_i - \overline{y})^2$		

Estimation and Inference

- The F-value is tested against a F-distribution with p, n-p-1 degrees of freedom
 - If we reject the null hypothesis, then the predictors do aid in predicting Y [in this analysis we do not know which ones are important!]
 - Failing to reject the null hypothesis does not mean that none of the covariates are important, since the effect of one or more covariates may be "masked" by others. The hard part is choosing which covariates to include or exclude.
- This is known as the global (multiple) F-test



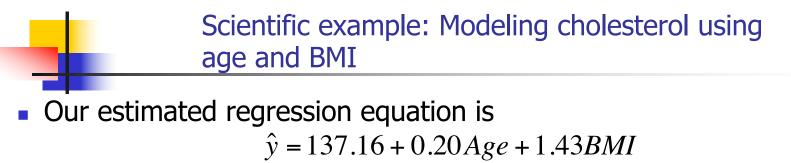
- We have seen that there is a significant relationship between age and cholesterol
- Can we better understand variability in cholesterol by incorporating additional covariates?





- It appears that BMI increases with increasing age
- And cholesterol increases with increasing BMI
- What if we want to estimate the association between age and cholesterol while holding BMI constant?
- Multiple regression!

```
> fit2=lm(chol~age+BMI)
> summary(fit2)
Call:
lm(formula = chol ~ age + BMI)
Residuals:
   Min
           1Q Median 3Q
                                  Max
-58.994 -15.793 0.571 14.159 62.992
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 137.1612
                       9.0061 15.230 < 2e-16 ***
            0.2023 0.0795 2.544 0.011327 *
age
             1.4266 0.3822 3.732 0.000217 ***
BMI
____
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 21.34 on 397 degrees of freedom
Multiple R-squared: 0.07351, Adjusted R-squared: 0.06884
F-statistic: 15.75 on 2 and 397 DF, p-value: 2.62e-07
```



• Question: How do we interpret the age coefficient?

- Our estimated regression equation is $\hat{y} = 137.16 + 0.20Age + 1.43BMI$
- Question: How do we interpret the age coefficient?
- Answer: This is the estimated average difference in cholesterol associated with a one year difference in age for two subjects with the same BMI.

Our estimated regression equation is

 $\hat{y} = 137.16 + 0.20 Age + 1.43 BMI$

- The age coefficient from our simple linear regression model was 0.31.
- Question: Why do the estimates from the two models differ?

Our estimated regression equation is

 $\hat{y} = 137.16 + 0.20 Age + 1.43 BMI$

- The age coefficient from our simple linear regression model was 0.31.
- Question: Why do the estimates from the two models differ?
- Answer: We are now conditioning on or controlling for BMI so our estimate of the age association is among subjects with the same BMI.



```
Call:
lm(formula = chol ~ age + BMI)
Residuals:
   Min
            10 Median
                           30
                                  Max
-58.994 -15.793 0.571 14.159 62.992
Coefficients:
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```

Cholesterol Example:

Did adding BMI improve our model?

```
> anova(fit,fit2)
Analysis of Variance Table
Model 1: chol ~ age
Model 2: chol ~ age + BMI
Res.Df RSS Df Sum of Sq F Pr(>F)
1 398 187187
2 397 1 80842 1 6345.8 13.931 0.0002174 ***
--- Signif. codes: 0 `***' 0.001 `**' 0.01 `*' 0.05 `.' 0.1 ` ' 1
```

How does the model with age and BMI compare to a model that contains only the mean?

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Interaction and Linear Regression

- Statistical interaction (aka effect modification) occurs when the relationship between an outcome variable and one predictor is different depending on the levels of a second predictor
- Interactions are usually investigated because of *a priori* assumptions/hypotheses on the part of the researchers
- Linear regression models allow for the inclusion of interactions with cross-product terms

Confounding vs. Interaction/Effect Modification

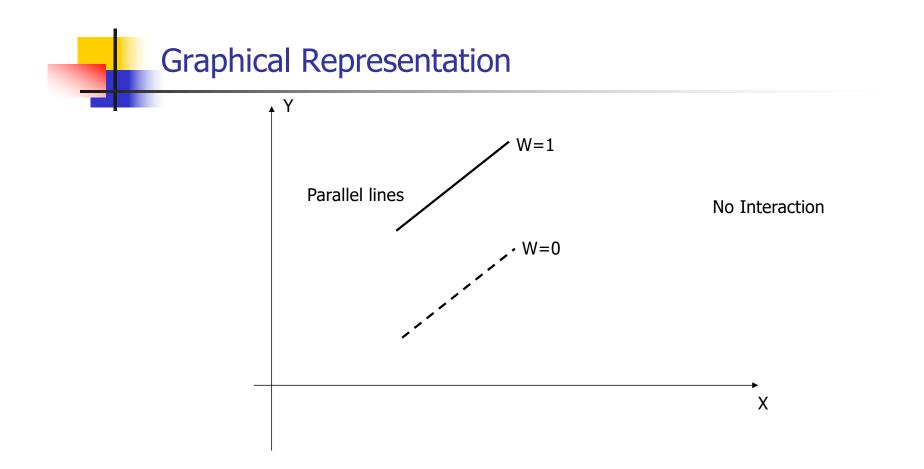
- Data and scientific understanding help distinguish between confounding and effect modifying variables:
 - Confounder: Associated with predictor and response; Association between response and predictor constant across strata of the new variable
 - Effect modifier/interaction: Association between response and the predictor varies across strata of the new variable

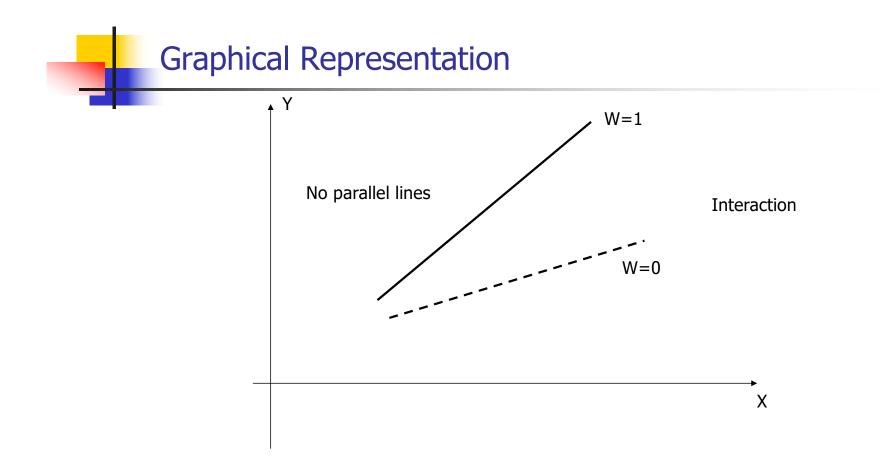
Confounding vs. Interaction/Effect Modification

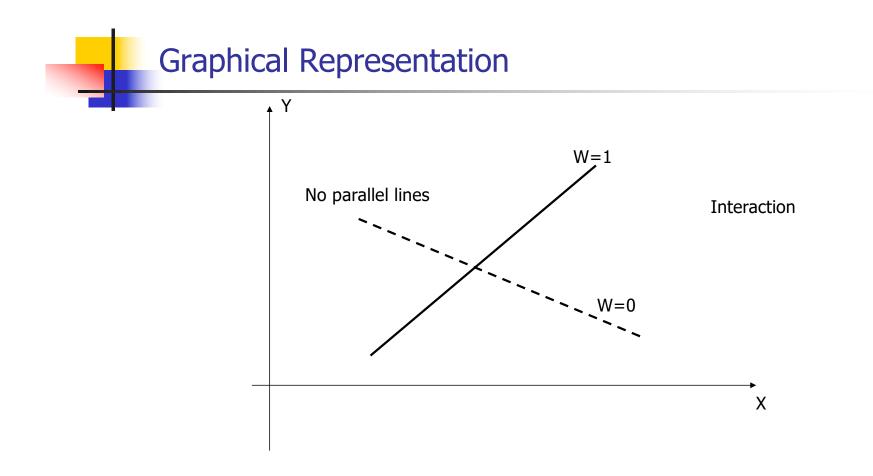
- Confounding: Estimates of association from unadjusted analysis are markedly different from estimates of association from adjusted analysis
 - Association within each stratum is similar, but different from the "crude" association in the combined data (ignoring the strata)
 - In linear regression, these symptoms are diagnostic of confounding
- Effect modification would show differences between adjusted analysis and unadjusted analysis, but would also show different associations in the different strata

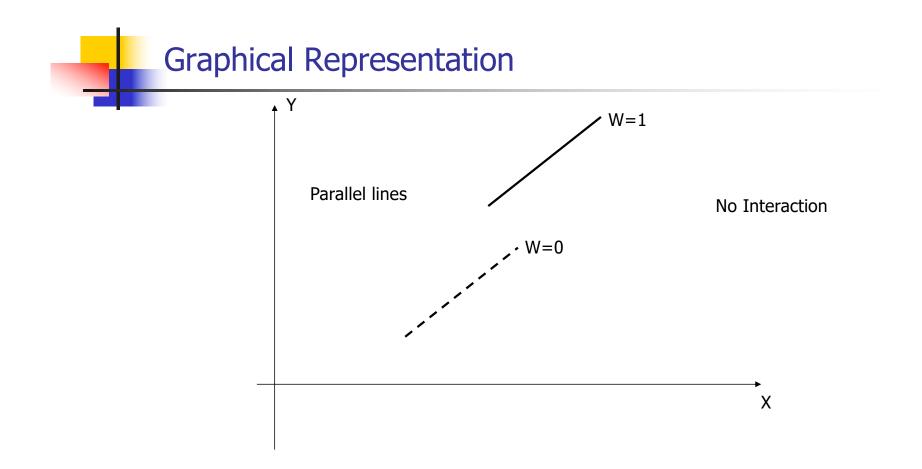
Effect Modification /Interaction

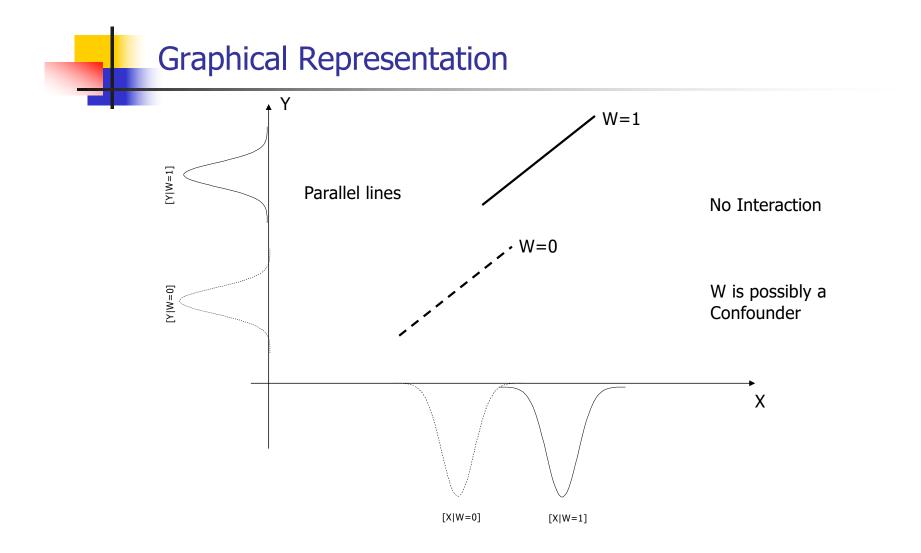
- Even if present, effect modification may not always be of interest in summarizing the effect of a predictor.
- For example, pleconaril, an antiviral drug, reduced the mean duration of symptoms in subjects with a common cold due to rhinoviruses but had no effect in subjects whose cold was due to some other agent.
- In the case of the pleconaril, effect modification was important in checking that the drug did actually work by inhibiting rhinovirus. However, in clinical use of the drug, it would typically not be possible to determine the infectious agent (the tests are expensive and take longer than just recovering from the cold), and so the average effectiveness of the drug across all colds would be a more important quantity.













Assume that there are two predictors in the model

 $\mathsf{E}[\mathsf{Y}|\mathsf{x}_{1},\mathsf{x}_{2}] = \beta_{0} + \beta_{1} \mathsf{x}_{1} + \beta_{2} \mathsf{x}_{2} + \beta_{3} \mathsf{x}_{1} \mathsf{x}_{2}$

Consider two observations with the same value, c, for x_2 , but one observation has x_1 one unit higher

Obs 1:
$$E[Y|x_1=k+1, x_2=c] = \beta_0 + \beta_1 (k+1) + \beta_2 c + \beta_3 (k+1)c$$

Obs 2: $E[Y|x_1=k, x_2=c] = \beta_0 + \beta_1 (k) + \beta_2 c + \beta_3 kc$

Thus, $E[Y|x_1=k+1, x_2=c] - E[Y|x_1=k, x_2=c] = \beta_1 + \beta_3 c$

That is, the difference in means depends now on the value of x_2 !

Model and Interpretation: interaction

- Model: $E[Y|x_1, x_2] = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_1 x_2$
- Difference in Means: $E[Y|x_1=k+1, x_2=c] - E[Y|x_1=k, x_2=c] = \beta_1 + \beta_3 c$

The difference in means depends on the value of x_2

- The difference in means is β_1 if c=0.
- The difference in means is $\beta_1 + \beta_3$ if c=1
- The difference in means changes by β_3 for each unit difference in c (that is, in x_2) [that is, β_3 is the difference of differences!]
- H₀: β₃=0 tests for interaction

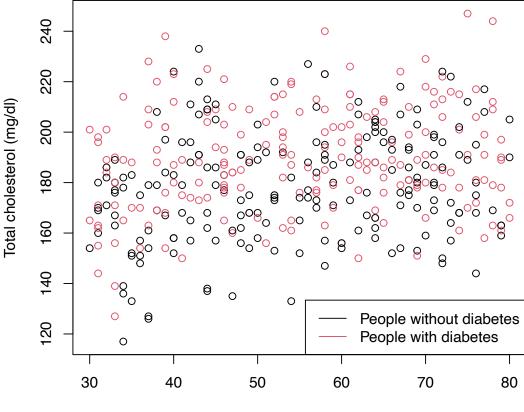
Model and Interpretation: interaction

- Model: $E[Y|x_1, x_2] = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_1 x_2$
- Another way to look at this
- Factor terms involving x₁: E[Y|x₁, x₂] = β₀+ (β₁ + β₃ x₂)x₁ + β₂ x₂

Slope of x₁ changes with x₂, i.e.

Difference in means for each unit difference in x_1 changes with x_2 (for each one unit difference in x_2 , the difference in means changes by β_3)

Cholesterol Example: Does diabetes affect the age – cholesterol relationship?



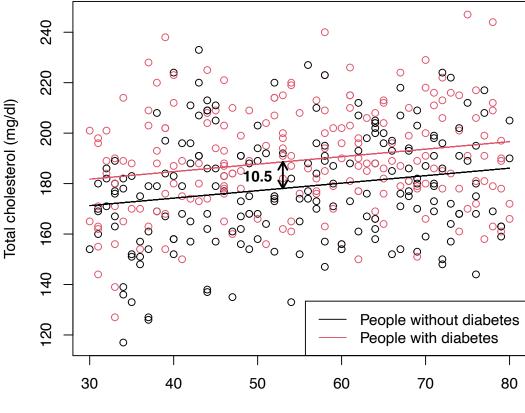
Age (years)



We first fit the model with age and DM terms only (No diabetes: DM=0, With diabetes: DM=1)

```
> fit3 = lm(chol ~ age+DM)
> summary(fit3)
Call:
lm(formula = chol ~ age + DM)
Residuals:
   Min
            10 Median
                           30
                                  Max
-55.662 -14.482 -1.411 14.682 57.876
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 162.35445 4.24184 38.275 < 2e-16 ***
             0.29697 0.07313 4.061 5.89e-05 ***
age
DM
            10.50728 2.10794 4.985 9.29e-07 ***
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 21.06 on 397 degrees of freedom
Multiple R-squared: 0.09748, Adjusted R-squared: 0.09293
F-statistic: 21.44 on 2 and 397 DF, p-value: 1.440e-09
```

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Age (years)

- This model indicates that, after controlling for the effect of diabetes, the average cholesterol differs by 0.30 for each additional year of age
- The age effect in this model is very similar to the effect from our simple linear regression (0.31)
- However, this does not mean that the age/cholesterol relationship is the same in people with and without diabetes
- To answer this question, we must add the interaction term



Model with age and DM main effects, plus interaction effect

```
> fit4=lm(chol~age*DM)
> summary(fit4)
Call:
lm(formula = chol ~ age * DM)
Residuals:
   Min
           10 Median 30
                                  Max
-56.474 -14.377 -1.215 14.764 58.301
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 160.31151 5.86268 27.344 < 2e-16 ***
age
            0.33460 0.10442 3.204 0.00146 **
           14.56271 8.29802 1.755 0.08004 .
DM
age:DM -0.07399 0.14642 -0.505 0.61361
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 21.08 on 396 degrees of freedom
Multiple R-squared: 0.09806, Adjusted R-squared: 0.09123
F-statistic: 14.35 on 3 and 396 DF, p-value: 6.795e-09
```

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Call: lm(formula :	= chol ~ age * DM)						
	1Q Median 3Q Max 377 -1.215 14.764 58.301						
Coefficient	3:						
	Estimate Std. Error t value Pr(> t)						
(Intercept)	160.31151 5.86268 27.344 < 2e-16 ***						
age	0.33460 0.10442 3.204 0.00146 **						
DM	14.56271 8.29802 1.755 0.08004 .						
age:DM	-0.07399 0.14642 -0.505 0.61361						
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1							
Residual standard error: 21.08 on 396 degrees of freedom							
Multiple R-squared: 0.09806, Adjusted R-squared: 0.09123 F-statistic: 14.35 on 3 and 396 DF, p-value: 6.795e-09							

Mean cholesterol for people without diabetes (DM = 0) at age 0



Call: lm(formula = chol ~ age * DM) Residuals: Min 10 Median 3Q Max -56.474 -14.377 -1.215 14.764 58.301 Coefficients: Estimate Std. Error t value Pr(>|t|) (Intercept) 160.31151 5.86268 27.344 < 2e-16 *** age 0.33460 0.10442 3.204 0.00146 ** DM 14.56271 8.29802 1.755 0.08004 . -0.07399 0.14642 -0.505 0.61361 age:DM ___ Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1 Residual standard error: 21.08 on 396 degrees of freedom Multiple R-squared: 0.09806, Adjusted R-squared: 0.09123 F-statistic: 14.35 on 3 and 396 DF, p-value: 6.795e-09

Difference in mean cholesterol between people with and without diabetes at age 0



Call: lm(formula =	= chol ~ age	* DM)					
	1Q Median .377 -1.215					/	
Coefficient	s:						
	Estimate St	d. Error	t value	Pr(> t)		/	
(Intercept)	160.31151	5.86268	27.344	< 2e-16	*** 🖌		
age	0.33460	0.10442	3.204	0.00146	* *		
DM	14.56271	8.29802	1.755	0.08004			
age:DM	-0.07399	0.14642	-0.505	0.61361			
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1							
Residual standard error: 21.08 on 396 degrees of freedom							
Multiple R-squared: 0.09806, Adjusted R-squared: 0.09123							
F-statistic: 14.35 on 3 and 396 DF, p-value: 6.795e-09							

Difference in mean cholesterol associated with each one-year change in age for people without diabetes



Call: lm(formula = chol ~ age * DM) Residuals: Min 10 Median 3Q Max -56.474 -14.377 -1.215 14.764 58.301 Coefficients: Estimate Std. Error t value Pr(>|t|) (Intercept) 160.31151 5.86268 27.344 < 2e-16 *** 0.10442 3.204 0.00146 ** age 0.33460 DM 14.56271 8.29802 1.755 0.08004 . 0.14642 -0.505 0.61361 age:DM -0.07399 ____ Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 Residual standard error: 21.08 on 396 degrees of freedom Multiple R-squared: 0.09806, Adjusted R-squared: 0.09123 F-statistic: 14.35 on 3 and 396 DF, p-value: 6.795e-09

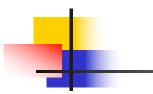
Difference in change in mean cholesterol associated with each one-year change in age comparing people with and without diabetes

- Interpretation?
 - Estimated model:

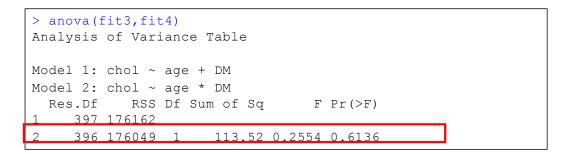
160.3 + 0.33 Age + 14.56 Diabetes - 0.07 Age x Diabetes

Subject 1: Age = a+1, diabetes = b Subject 2: Age = a, diabetes = b Difference in the estimated cholesterol: [160.3 + 0.33(a+1) + 14.56(b) - 0.07 (a+1)(b)] -[160.3 + 0.33(a) + 14.56 (b) - 0.07 (a)(b)] = 0.33-0.07b

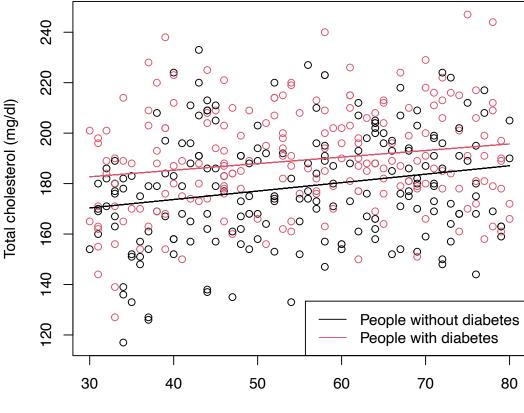
 Diabetes exerts a small (not statistically significant) effect on the age/cholesterol relationship In people without diabetes: 160.3+0.33 Age In people with diabetes : 174.9+0.26 Age



 We can also test the significance of interaction terms using an F-test



 Adding the interaction term did not significantly improve model fit



Age (years)



We have considered:

- Simple linear regression
 - Interpretation
 - Estimation
 - Model checking
- Multiple linear regression
 - Confounding
 - Interpretation
 - Estimation
 - Interaction



• Let's work on Exercises 4-8