

HIP 529 Applied Biostatistics II

DATE: March 28, 2022

PRESENTED BY: Emile Latour, MS, Associate Biostatistician, OHSU Knight Cancer Institute

Slides are available

https://bit.ly/hip-529-logistic-regression_2022-03-28



About me

- Pension actuary for 16 years
- MS in Biostatistics from OHSU in 2017
- Associate Biostatistician with OHSU Knight Cancer Institute since 2016
- Long term support for OHSU Dermatology Department since 2018





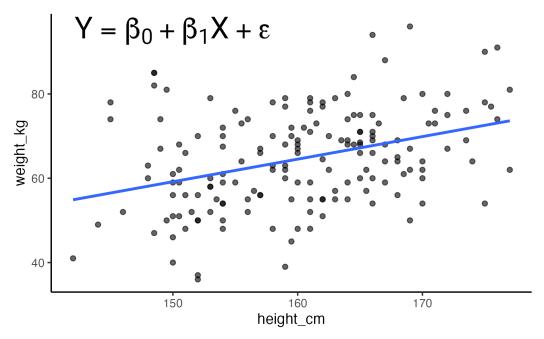
Goals

- Exposure to statistics background on logistic regression models and what sets it apart
- Understand terms: log odds, odds ratio, and relative risk
- Univariable logistic regression models (association models):
 - Continuous predictor, categorical predictors



Assumed prerequisite

Some familiarity with simple linear regression





Assumed prerequisite

- Linear regression the goal is to come up with a linear equation that describes the relationship between dependent and independent variables.
- Logistic regression the goal is to develop an equation that describes the probability of belonging to one group or another.





Goal of statistical modeling

• To find the best fitting and most parsimonious, yet biologically reasonable model to describe the relationship between an outcome variable and a set of independent variables.



Describe a relationship

 A statistical model is a mathematical relationship between an outcome (dependent variable) and one or more independent variables.



Describe a relationship

 A statistical model is a mathematical relationship between an outcome (dependent variable) and one or more independent variables.





Types of data / variables

- Discrete (numerical) quantities or counts, likely with units
- Continuous (numerical) -- quantities, likely with units
- Ordinal (categorical) labels with order
- Nominal (categorical) labels with no order



Discrete (numerical) data

- Actual quantities; not numerical coding of categories
- Restricted to a set of isolated points
- Intermediate values are impossible
- Examples:
 - Occupants per household: 1, 2, 3,
 - Number of tumors: 0, 1, 2, 3, 4, ...
 - Age to nearest whole year: ..., 35, 36, 37, ...



Continuous (numerical) data

- Actual quantities
- Not restricted to a set of isolated points
- Intermediate values are(theoretically) possible
- Examples:
 - Change in body weight during pandemic: +19.25 lbs
 - Cholesterol level: 162 mg/dL
 - Age (fractional): 35.75 years



Ordinal (categorical) data

- Categories or classes with meaningful order
- Examples:
 - Pain: None, Mild, Moderate, Severe
 - Likert: Disagree, Neutral, Agree
 - Smoking status: Never, Occasionally, Frequently
- Numerical codes can be assigned, but must show the ranking (order)



Nominal (categorical) data

- Unordered categories or classes
- Examples:
 - Biological sex: M = Male, F = Female
 - Blood type: A, B, O, AB
 - Smoking status: Yes/No (1/0)
- Data can be given numerical codes (e.g. 1=A, 2=B, 3=O, 4=AB)



Special case – Binary data

- Nominal data with only two levels
- Examples: Male/Female, Yes/No, Case/Control, Tx/No Tx
- Labels can be replaced with the codes 0 and 1
 - Sum of the numeric codes gives the total with code 1
 - The average of the codes gives the proportion (or percentage) assigned the code 1
 - Example: Sample of n = 10 people coded as smoker (1) or non-smoker (0):
 - {1, 0, 1, 0, 1, 1, 1, 1, 1, 0}
 - Total = 7
 - Average = Total / n = 7 / 10 = 0.70 = 70.0%





Regression analysis

 Mathematically describe the relationship between a dependent variable (outcome) and a set of one or more independent variables



Choosing a model

 Most often, the type of data of your outcome will determine the type of model that is used



Terms

- Dependent variable, aka
 - Outcome
 - Response
 - Typically denoted with Y
- Independent variable(s)
 - Predictors
 - Explanatory variables
 - Typically denoted with X_i



Classes of models

- "General" Linear models:
 - Refers to conventional linear regression models
 - Continuous outcome variable
 - Continuous and categorical predictors
 - Also includes multiple linear regression, ANOVA, and ANCOVA



Classes of models

- "Generalized" Linear models (GLM):
 - Broad class of models
 - Includes ordinary regression/ANOVA models for continuous outcomes
 - Also, includes models that can handle categorical outcomes



3 components of a GLM

- Random component
- Systematic component
- Link function



Random component

- Specifies the response variable Y and a probability distribution for it
 - If Y is continuous, then assumes normal distribution of Y
 - If Y is binary, then assumes a binomial distribution of Y
 - If Y is a count, then assumes a Poisson or negative binomial distribution of Y



Binomial distribution

- Success/Failure
- Conditions:
 - The number of trials is fixed (n)
 - Each trial has one of two outcomes: success or failure
 - The probability of success (p) is the same for each trial
 - Trials are independent; the outcome of one trial does not affect the outcome of another



Binomial distribution

- Examples
 - Number of heads out of 20 coin flips
 - Number of times you hit red light on your commute to work
 - Number of passengers on the Titanic that survived
 - Number of patients with a disease in a sample of 1000 patients.



Systematic component

• Specifies the explanatory variables $(x_1, x_2, ..., x_k)$ which enter the model as a linear combination

$$\beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_k x_k$$

- Here the x_i can be based on other explanatory variables:
 - For example, $x_3 = x_1 * x_2$ or $x_3 = x_1^2$



Link function

- Connects the predictors in a model with the expected value of the response variable in a linear way
- Specifies the link between the random component and the systematic component



Link function

Let the expected value of the response be

$$\mu = E(Y)$$

• The link function specifies g(.) that relates μ to the linear combination of predictors

$$g(\mu) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_k x_k$$



Link function

- Identity link
 - The link for a continuous response variable
 - $-g(\mu)=\mu$
- Log link
 - Often used for a discrete count response variable
 - $g(\mu) = log(\mu)$
- Logit link
 - Often used for a categorical response variable
 - $g(\mu) = logit(\mu) = log\left(\frac{\mu}{1-\mu}\right)$
 - A GLM that uses a logit link function is called a logistic regression model



GLM for binary data

- Identity link is not appropriate when the dependent variable is categorical
- Assumptions that underlie ordinary linear regression (OLR) are violated:
 - Linear relationship between μ and x can be problematic
 - Normality assumption is violated
 - Serious when sample size is small; less serious when sample size is large. Still an issue either way
 - Homoscedasticity assumption is violated
 - Can lead to biased estimates of the standard error to an unknown degree





CHD Study

 A study to evaluate the relationship between age and presence of coronary heart disease (CHD)



CHD Study

- N = 100
- First 20 shown here

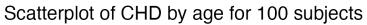
id	age	agegrp	chd
1	20	20-29	No
2	23	20-29	No
3	24	20-29	No
4	25	20-29	No
5	25	20-29	Yes
6	26	20-29	No
7	26	20-29	No
8	28	20-29	No
9	28	20-29	No
10	29	20-29	No
11	30	30-34	No
12	30	30-34	No
13	30	30-34	No
14	30	30-34	No
15	30	30-34	No
16	30	30-34	Yes
17	32	30-34	No
18	32	30-34	No
19	33	30-34	No
20	33	30-34	No

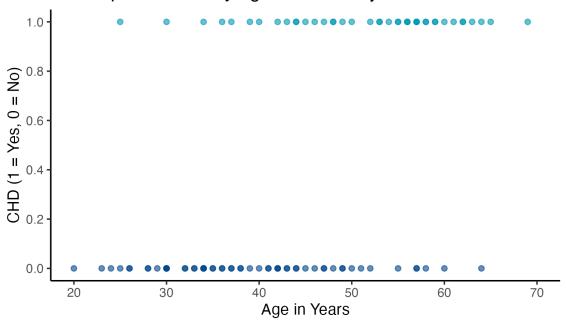


CHD Study

- Primary endpoint (dependent variable): CHD
 - CHD = "Yes": patient has coronary heart disease
 - CHD = "No": patient does not have coronary heart disease



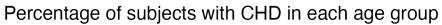


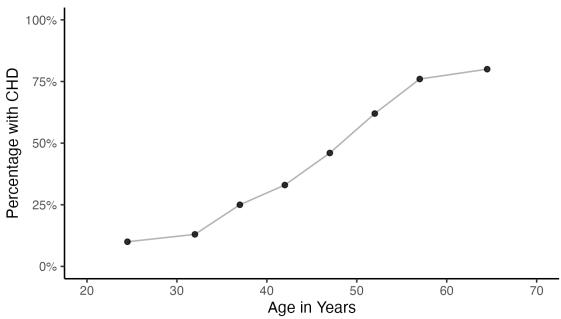




Frequency table of CHD by age group								
	CHD							
Age group	n	Yes	No	Mean (proportion)				
20-29	10	1	9	0.10				
30-34	15	2	13	0.13				
35-39	12	3	9	0.25				
40-44	15	5	10	0.33				
45-49	13	6	7	0.46				
50-54	8	5	3	0.62				
55-59	17	13	4	0.76				
60-69	10	8	2	0.80				
Total	100	43	57	0.43				







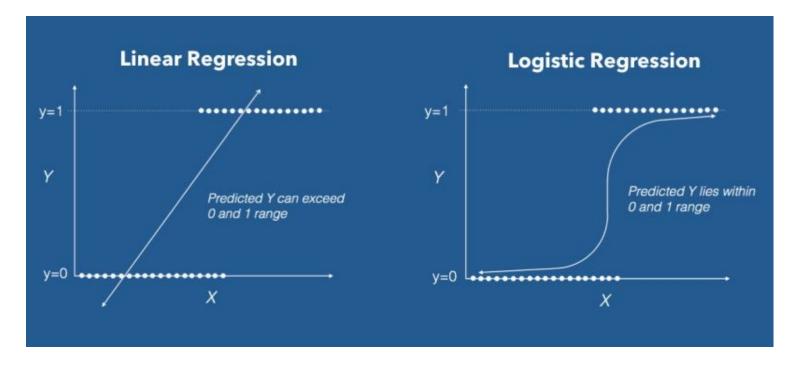


- Apparent trend: increasing proportion with CHD as age increases
- Relationship does not look linear in the two ends





Another look at the issue





Issue

- We have binary data
- We want to be able to run something like a regression
- But where we model the probability of the outcome



Issue and solution

- Probabilities are limited between 0 and 1
- To use a linear model we need to transform the probabilities so that they range from $-\infty$ (infinity) to $+\infty$



- Transform the probabilities
 - From 0 to 1
 - To 0 to ∞
- Can do this by converting probabilities to odds, expressed as a ratio

$$odds = \frac{p}{1 - p}$$



 Probabilities and odds are two equivalent ways of expressing the same idea

$$odds = \frac{p}{1 - p}$$

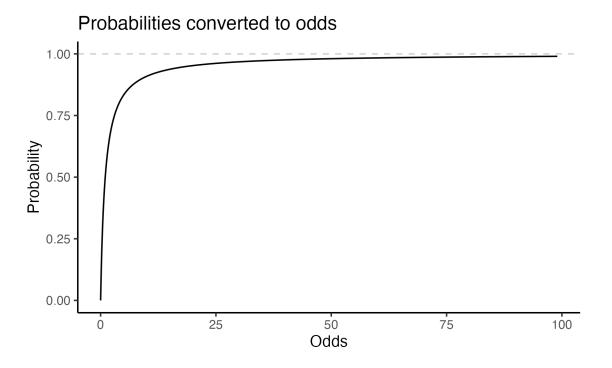
$$p = \frac{odds}{1 + odds}$$



• Say
$$p = 0.5$$
 then $odds = \frac{0.5}{1 - 0.5} = \frac{0.5}{0.5} = \frac{1}{1}$ or 1 to 1

• Say
$$p = 0.25$$
 then $odds = \frac{0.25}{1 - 0.25} = \frac{0.25}{0.75} = \frac{1}{3}$ or 1 to 3







- If we convert probability to odds, then the odds are always > zero
- For a linear model, we want our estimates to range from $-\infty$ to $+\infty$



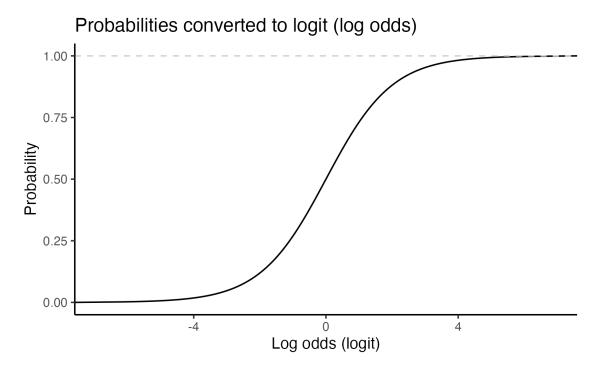
Log odds

- To solve this, we can take the logarithm of the odds to get our estimates on a scale $-\infty$ to $+\infty$
- Logarithm of the odds is called the logit

$$logit(p) = log(odds) = log\left(\frac{p}{1-p}\right)$$



Logit





Logit

• By taking the log odds (logit) of the probability, our estimates on a scale $-\infty$ to $+\infty$



Logistic regression model

$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_k x_k$$

- Random component follows a binomial distribution
- Systematic component
- Link function: logit link



Logistic regression model

- Units are in log-odds
- In linear regression, a coefficient (β) is the change in the outcome for a unit change in the predictor.
- In logistic regression, a coefficient is the change in log-odds of the outcome being 1 for a unit change in the predictor.



Logistic regression model

- Change in log-odds is the same as the ratio of odds.
- To get an odds ratio, need to transform the log-odds.
- Exponentiate the estimated model coefficient

$$Odds \ ratio = \exp(\beta) = e^{\beta}$$





Hip fracture study

• A study to examine the relationship between hip fracture with age, sex, BMI, and bone mineral density (bone densitometries).



Hip fracture study

- id: subject ID
- age: subject age in years
- sex: biological sex (Female/Male)
- fracture: hip fracture (Fracture/Nofracture)
- weight_kg: weight in kg
- height_cm: height in cm
- medication: prescribed medications
- waiting_time: time the subject waited for the densitometry (in minutes)
- bmd: bone mineral density measure in the hip
- bmi: calculated body mass index
- bmi_c: Body mass index category (Underweight, Healthy, Overweight, Obese)



https://bookdown.org/tpinto_home/Regression-and-Classification/#datasets-used-in-the-examples

• N = 169; First 20 shown below

id	age	sex	fracture	weight_kg	height_cm	medication	waiting_time	bmd	bmi	bmi_c
00469	57.05	Female	No	64	155.5	Anticonvulsant	18	0.8793	26.46788	Overweight
08724	75.74	Female	No	78	162.0	No medication	56	0.7946	29.72108	Overweight
06736	70.78	Male	No	73	170.5	No medication	10	0.9067	25.11158	Overweight
24180	78.25	Female	No	60	148.0	No medication	14	0.7112	27.39226	Overweight
17072	54.19	Male	No	55	161.0	No medication	20	0.7909	21.21832	Healthy
03806	77.18	Male	No	65	168.0	No medication	7	0.7301	23.03005	Healthy
17106	56.18	Male	No	77	159.0	No medication	26	1.0096	30.45766	Obese
23834	49.92	Female	No	59	150.0	No medication	9	0.7310	26.22222	Overweight
02454	68.41	Male	No	64	167.0	Glucocorticoids	6	0.6893	22.94812	Healthy
02088	66.26	Male	No	72	159.5	No medication	10	0.9466	28.30161	Overweight
05364	45.87	Male	No	62	169.0	No medication	12	0.8015	21.70792	Healthy
08922	73.97	Female	No	68	164.0	No medication	5	0.5793	25.28257	Overweight
23890	60.56	Female	No	76	155.0	No medication	11	0.9760	31.63371	Obese
03047	64.21	Male	No	90	175.0	Glucocorticoids	28	0.9184	29.38776	Overweight
02179	53.40	Male	No	70	162.5	No medication	73	0.8020	26.50888	Overweight
03800	66.83	Male	No	76	171.0	No medication	13	0.8033	25.99090	Overweight
07528	57.93	Male	No	67	160.0	Glucocorticoids	5	0.7978	26.17187	Overweight
05288	40.23	Male	No	66	165.0	No medication	8	1.0390	24.24242	Healthy
00109	69.05	Female	No	72	154.0	No medication	11	0.7861	30.35925	Obese
08622	57.80	Female	No	50	152.0	No medication	21	0.8254	21.64127	Healthy



Hip fracture Study

- Primary endpoint (dependent variable): fracture
 - fracture = "Yes": patient had a hip fracture
 - fracture = "No": patient did not have hip fracture



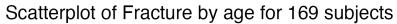
- Research question #1:
 - Is there a relationship between a hip fracture occurring and age?

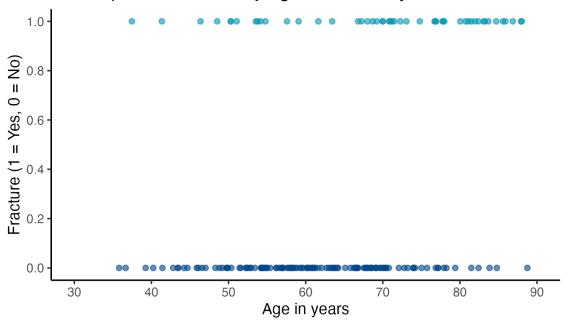


fracture	n	percent
Yes	50	29.6%
No	119	70.4%
Total	169	100.0%

fracture	variable	n	complete	missing	mean	sd	p0	p25	p50	p75	p100	range
All subjects	age	169	169	0	63.63	12.36	35.81	54.42	63.49	72.08	88.75	52.94
Yes	age	50	50	0	69.77	13.38	37.46	59.72	71.32	81.01	88.02	50.56
No	age	119	119	0	61.05	10.97	35.81	54.21	60.56	68.56	88.75	52.94









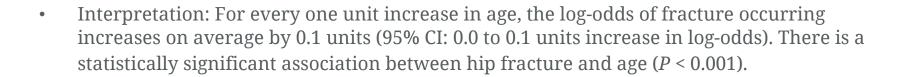
$$logit(p) = \beta_0 + \beta_1 Age$$

- p = probability of Fracture
- β_1 determines whether the curve ascends or descends
 - $-\beta_1 > 0$, then *p* increases as *Age* increases
 - $-\beta_1 < 0$, then p decreases as Age increases
 - $-\beta_1 = 0$, then p is independent of Age



Fracture and age – log-odds

term	estimate	lower_ci	upper_ci	p_value
(Intercept)	-5.0	-7.2	-3.0	< 0.001
age	0.1	0.0	0.1	< 0.001





Fracture and age – log-odds

```
> res <- glm((fracture == "Yes") ~ age,
+ family = binomial(link = "logit"),
+ data = data)
> summary(res)
Call:
glm(formula = (fracture == "Yes") ~ age, family = binomial(link = "logit"),
   data = data)
Deviance Residuals:
   Min 1Q Median 3Q Max
-1.4444 -0.8553 -0.6216 1.0699 2.3290
Coefficients:
        Estimate Std. Error z value Pr(>|z|)
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 205.27 on 168 degrees of freedom
Residual deviance: 186.76 on 167 degrees of freedom
AIC: 190.76
Number of Fisher Scoring iterations: 4
```



Fracture and age – log-odds

```
. logit fracture age
Iteration 0: log likelihood = -102.63604
Iteration 1: log likelihood = -93.602157
Iteration 2: log likelihood = -93.378565
Iteration 3: log likelihood = -93.37797
Iteration 4: log likelihood = -93.37797
Logistic regression
                                LR chi2(1)
                                                 18.52
                                Prob > chi2
                                                 0.0000
Log likelihood = -93.37797
                                Pseudo R2
______
  fracture | Coef. Std. Err. z P>|z| [95% Conf. Interval]
     age | .0634076 .0158255 4.01 0.000 .0323902 .094425
```



Fracture and age – odds ratio

$$logit(p) = \beta_0 + \beta_1 Age$$

$$Odds \ ratio = OR = e^{\beta_1}$$

- OR > 1, then greater likelihood of outcome (hip fracture) as age increases
- OR < 1, then lesser likelihood of outcome as age increases
- -OR = 1, then the same likelihood of outcome as age increases



Fracture and age -odds ratio

term	estimate	lower_ci	upper_ci	p_value	
(Intercept)	0.0	0.0	0.0	< 0.001	
age	1.1	1.0	1.1	< 0.001	

• Interpretation: For every one unit increase in age, the odds of fracture occurring increases 1.1 times (95% CI: 1.0 to 1.1 times). There is a statistically significant association between hip fracture and age (P < 0.001).



Fracture and age -odds ratio



Fracture and age -odds ratio

```
. logit fracture age, or
Iteration 0: log likelihood = -102.63604
Iteration 1: log likelihood = -93.602157
Iteration 2: log likelihood = -93.378565
Iteration 3: log likelihood = -93.37797
Iteration 4: log likelihood = -93.37797
Logistic regression
                                     LR chi2(1)
                                                         18.52
                                     Prob > chi2
                                                        0.0000
Log likelihood = -93.37797
                                     Pseudo R2
______
   fracture | Odds Ratio Std. Err. z P>|z| [95% Conf. Interval]
      age | 1.065461 .0168614 4.01 0.000
     _cons | .0066136 .007121 -4.66 0.000 .0008016 .0545688
```



Fracture and sex

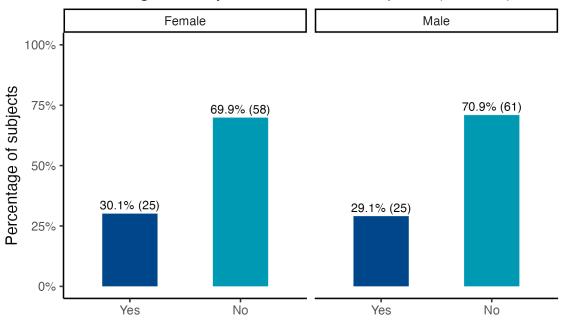
- Research question #2:
 - Is there a relationship between a hip fracture occurring and biological sex?



fracture/sex	Male	Female	Total
Yes	25	25	50
	(29.1%)	(30.1%)	(29.6%)
No	61	58	119
	(70.9%)	(69.9%)	(70.4%)



Percentage of subjects with Fracture by sex (n = 169)





$$logit(p) = \beta_0 + \beta_1 Sex$$

- p = probability of Fracture
- β_1 determines whether the curve ascends or descends
 - $-\beta_1 > 0$, then *p* higher in males than females
 - $-\beta_1 < 0$, then *p* lower in males than females
 - $-\beta_1 = 0$, then p the same in males than females



Fracture and sex – odds ratio

$$logit(p) = \beta_0 + \beta_1 sex$$

$$Odds \ ratio = OR = e^{\beta_1}$$

- OR > 1, then greater likelihood of outcome (hip fracture) as in males than females
- OR < 1, then lesser likelihood of outcome in males than females
- OR = 1, then the same likelihood of outcome in males than females



term	level	reference	estimate	lower_ci	upper_ci	p_value
(Intercept)	(Intercept)	Non- Baseline Category	0.4	0.3	0.7	< 0.001
sex	Female	Baseline Category				
sex	Male	Non- Baseline Category	1.0	0.5	1.8	0.881

• Interpretation: The odds of hip fracture among male subjects is 1.0 times (95% CI: 0.5 to 1.8 times) the odds of hip fracture among female subjects. There is no statistically significant association between hip fracture and sex (P < 0.881).



```
> res <- glm((fracture == "Yes") ~ sex,
          family = binomial(link = "logit"),
           data = data)
> summary(res)
glm(formula = (fracture == "Yes") ~ sex, family = binomial(link = "logit"),
   data = data)
Deviance Residuals:
   Min 1Q Median 3Q Max
-0.8466 -0.8466 -0.8288 1.5492 1.5719
Coefficients:
          Estimate Std. Error z value Pr(>|z|)
sexMale -0.05043 0.33710 -0.150 0.881078
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 205.27 on 168 degrees of freedom
Residual deviance: 205.25 on 167 degrees of freedom
AIC: 209.25
Number of Fisher Scoring iterations: 4
> ## odds ratios and 95% CI
> exp(cbind(OR = coef(res), confint(res)))
Waiting for profiling to be done...
                OR 2.5 % 97.5 %
(Intercept) 0.4310345 0.2651600 0.6804247
sexMale 0.9508197 0.4898953 1.8448256
```



```
. logit fracture i.sex, or
Iteration 0: log likelihood = -102.63604
Iteration 1: log likelihood = -102.62485
Iteration 2: log likelihood = -102.62485
Logistic regression
                                    Number of obs
                                                         169
                                    LR chi2(1)
                                                        0.02
                                    Prob > chi2
Log likelihood = -102.62485
                                    Pseudo R2
   fracture | Odds Ratio Std. Err. z P>|z| [95% Conf. Interval]
-----
      sex
     Male | .9508197 .3205191 -0.15 0.881 .4910889 1.840925
     _cons | .4310345 .1031257 -3.52 0.000 .2696874 .6889115
```



Reference levels

- For interpretation it's vital to know which level of a variable is the Reference Level.
- The level of a categorical variable considered the "baseline" or "usual" value
- Other levels are compared to the Reference Level



Reference levels

- In the Fracture and Sex example, *Female* is the reference level.
- The models give us an odds ratio for the *Male* level.
- Odds ratio is in terms of Male compared to Female.



Reference levels

- When in doubt, check it.
- Can do this "by hand" with a 2x2 table.



fracture/sex	Male	Female	Total
Yes	25	25	50
	(29.1%)	(30.1%)	(29.6%)
No	61	58	119
	(70.9%)	(69.9%)	(70.4%)

Yes Outcome No

Yes	No
a	b
с	d

Predictor



•
$$a = 25$$

•
$$b = 25$$

•
$$c = 61$$

•
$$d = 58$$

fracture/sex	Male	Female	Total
Yes	25	25	50
	(29.1%)	(30.1%)	(29.6%)
No	61	58	119
	(70.9%)	(69.9%)	(70.4%)



•
$$a = 25$$

•
$$b = 25$$

•
$$c = 61$$

•
$$d = 58$$

fracture/sex	Male	Female	Total
Yes	25	25	50
	(29.1%)	(30.1%)	(29.6%)
No	61	58	119
	(70.9%)	(69.9%)	(70.4%)

• $Odds\ ratio = OR = (a/b)/(c/d) = ad/bc$

•
$$OR = (25/25)/(61/58) = (25 * 58)/(25 * 61) = 0.95$$



Predictor

 Yes
 No

 Yes
 b

 Outcome
 No
 c
 d

• The odds of [outcome = "Yes"] among those [predictor = "Yes"] is [odds ratio] times the odds of those with [predictor = "No].



sex/fracture	Yes	No	Total	
Male	25	61	86	
	(50.0%)	(51.3%)	(50.9%)	
Female	25	58	83	
	(50.0%)	(48.7%)	(49.1%)	

Yes Predictor No

Yes	No
a	b
с	d

Outcome



•
$$a = 25$$

•
$$b = 61$$

•
$$c = 25$$

•
$$d = 58$$

sex/fracture	Yes	No	Total	
Male	25	61	86	
	(50.0%)	(51.3%)	(50.9%)	
Female	25	58	83	
	(50.0%)	(48.7%)	(49.1%)	

•
$$OR = (25/61)/(25/58) = (25 * 58)/(61 * 25) = 0.95$$

Same result as before



Change the reference level – 2x2 table

•	а	=	25
	а		

•
$$b = 25$$

•
$$c = 58$$

•
$$d = 61$$

fracture/sex	Female	Male	Total
Yes	25	25	50
	(30.1%)	(29.1%)	(29.6%)
No	58	61	119
	(69.9%)	(70.9%)	(70.4%)

•
$$Odds \ ratio = OR = (a/b)/(c/d) = ad/bc$$

•
$$OR = (25/25)/(58/61) = (25 * 61)/(25 * 58) = 1.05$$

• Different than before! Pay attention to the reference level.



Change the reference level – LR Model

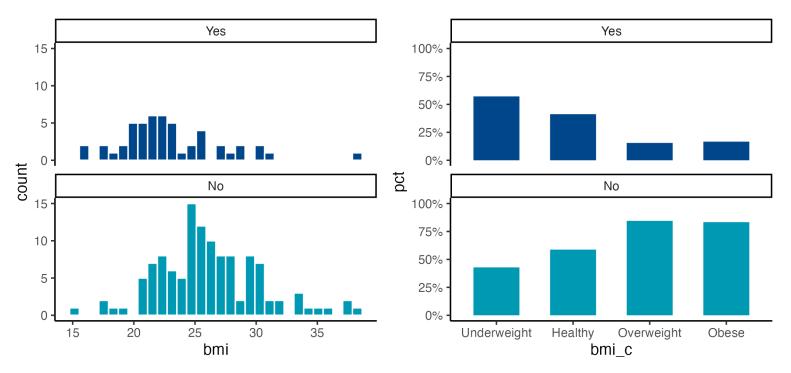
term	level	reference	estimate	lower_ci	upper_ci	p_value
(Intercept)	(Intercept)	Non- Baseline Category	0.4	0.3	0.6	< 0.001
sex	Male	Baseline Category				
sex	Female	Non- Baseline Category	1.1	0.5	2.0	0.881



- Research question #2:
 - Is there a relationship between a hip fracture occurring and BMI?



BMI – continuous or categorical?





- Categorical BMI
- "Healthy" is the reference level



term	level	reference	estimate	lower_ci	upper_ci	p_value
(Intercept)	(Intercept)	Non- Baseline Category	0.7	0.4	1.1	0.119
bmi_c	Underweight	Non- Baseline Category	1.9	0.4	10.2	0.421
bmi_c	Healthy	Baseline Category				
bmi_c	Overweight	Non- Baseline Category	0.3	0.1	0.6	0.002
bmi_c	Obese	Non- Baseline Category	0.3	0.1	0.8	0.034



- Interpret relative to the Reference Level
 - Compared to Healthy subjects, Underweight subjects were 1.9 times (95% CI: 0.4 to 10.2 times) more likely to have hip fracture (P = 0.421).
 - The odds of hip fracture among Overweight subjects was 0.3 times (95% CI: 0.1 to 0.6 times) the odds of hip fracture among Healthy subjects. Overweight subjects were less likely to experience hip fracture (P = 0.002).



```
> data$bmi c <- relevel(data$bmi c, ref = "Healthy")</pre>
> res <- glm((fracture == "Yes") ~ bmi_c,</pre>
           family = binomial(link = "logit"),
           data = data)
> summary(res)
Call:
glm(formula = (fracture == "Yes") ~ bmi_c, family = binomial(link = "logit"),
   data = data)
Deviance Residuals:
   Min 10 Median 30
-1.3018 -1.0314 -0.5807 1.3308 1.9304
Coefficients:
                Estimate Std. Error z value Pr(>|z|)
(Intercept)
                -0.3536 0.2271 -1.557 0.11944
bmi_cUnderweight 0.6413
                           0.7968 0.805 0.42090
                           0.4279 -3.134 0.00173 **
bmi_cOverweight -1.3410
bmi_cObese
                 -1.2558
                           0.5929 -2.118 0.03418 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 205.27 on 168 degrees of freedom
Residual deviance: 189.69 on 165 degrees of freedom
AIC: 197.69
Number of Fisher Scoring iterations: 4
> ## odds ratios and 95% CI
> exp(cbind(OR = coef(res), confint(res)))
Waiting for profiling to be done...
                      OR 2.5 % 97.5 %
               0.7021277 0.44643887 1.0914867
bmi cUnderweight 1.8989899 0.39391648 10.1755229
bmi_cOverweight 0.2615955 0.10772680 0.5853194
```

0.2848485 0.07743896 0.8363863

bmi cObese



```
. logit fracture ib1.bmi_c2, or
Iteration 0: log likelihood = -102.63604
Iteration 1: log likelihood = -95.007409
Iteration 2: log likelihood = -94.846001
Iteration 3: log likelihood = -94.845749
Iteration 4: log likelihood = -94.845749
Logistic regression
                                      LR chi2(3)
                                      Prob > chi2
                                                          0.0014
Log likelihood = -94.845749
                                      Pseudo R2
   fracture | Odds Ratio Std. Err. z P>|z| [95% Conf. Interval]
-----
    bmi_c2 |
    Obese .2848485 .1688985 -2.12 0.034
                                              .0891048
                                                        .9105979
Overweight | .2615955 .1119369 -3.13 0.002
                                              .1130839
                                                        .6051457
Underweight |
             1.89899 1.513142
                                              .3983576
                                                        9.052576
     _cons | .7021277 .1594613 -1.56 0.119
                                              .4498818 1.095806
```



Odds and ends

- Relative Risk
 - In case control studies and cohort studies where the outcome occurs in less than 10% of the exposed population, Odds Ratio provides a reasonable approximation to the Relative Risk



Odds and ends

- Association vs. Prediction
 - Models we have seen here are focused on associations – relationships and interpretations
 - Prediction/classification aims to predict. Topic for another time



Odds and ends

- Univariable one explanatory variable
- Univariate One response variable
- Multivariable more than one explanatory variable
- Multivariate more than one response variable



References

- An introduction to logistic regression: from basic concepts to interpretation with particular attention to nursing domain (Park 2013)
- <u>Understanding logistic regression analysis (Sperandei)</u>
- <u>Using logistic regression in perinatal epidemiology: an introduction for clinical researchers. Part 1: basic concepts (Brand, Keirse)</u>
- An introduction to logistic regression with an application to the analysis of language recovery following a stroke (Greenhouse et al)
- Explaining Odds Ratios (Szumilas)
- Penn State, STAT 504, Binary Logistic Regression
- Developing prediction models for clinical use using logistic regression: an overview (Shipe et al)





Thank You