

432 Class 8 Slides

github.com/THOMASELOVE/432-2018

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Setup

```
library(skimr)  
library(broom)  
library(Hmisc)  
library(rms)  
library(tidyverse)
```

Today's Materials

- Logistic Regression and the Low Birth Weight data

```
lbw1 <- read.csv("data/lbw.csv") %>% tbl_df

lbw1 <- lbw1 %>%
  mutate(race_f = fct_recode(factor(race), white = "1",
                                   black = "2", other = "3"),
         race_f = fct_relevel(race_f, "white", "black")) %>%
  mutate(preterm = fct_recode(factor(ptl > 0),
                                   yes = "TRUE",
                                   no = "FALSE")) %>%
  select(subject, low, lwt, age, ftv, ht, race_f,
         preterm, smoke, ui)
```

The lbw1 data (n = 189 infants)

Variable	Description
subject	id code
low	indicator of low birth weight (< 2500 g)
lwt	mom's weight at last menstrual period (lbs.)
age	age of mother in years
ftv	count of physician visits in first trimester (0 to 6)
ht	history of hypertension: 1 = yes, 0 = no
race_f	race of mom: white, black, other
preterm	prior premature labor: 1 = yes, 0 = no
smoke	1 = smoked during pregnancy, 0 = did not
ui	presence of uterine irritability: 1 = yes, 0 = no

Source: Hosmer, Lemeshow and Sturdivant, *Applied Logistic Regression* 3rd edition. Data from Baystate Medical Center, Springfield MA in 1986.

Goals for Today and Tuesday

- 1 Fit and evaluate the fit of a logistic regression model to predict the probability of a low birth weight ($\text{low} = 1$) using the mom's weight at her last menstrual period (lwt).
- 2 Fit and evaluate a larger logistic regression model to predict low on the basis of a larger group of predictors drawn from the available options, which include: lwt , age , ftv , ht , race_f , preterm , smoke and ui .
- 3 Learn about the use of both glm and lrm (from the rms package) to fit and evaluate logistic regression models.

EDA for Task 1

We want to look at the probability of a low birth weight ($\text{low} = 1$) on the basis of the mom's weight at her last menstrual period (lwt).

```
lbw1 %>% group_by(low) %>% skim(lwt)
```

```
> lbw1 %>% group_by(low) %>% skim(lwt)
```



```
Skim summary statistics
```

```
n obs: 189
```

```
n variables: 10
```

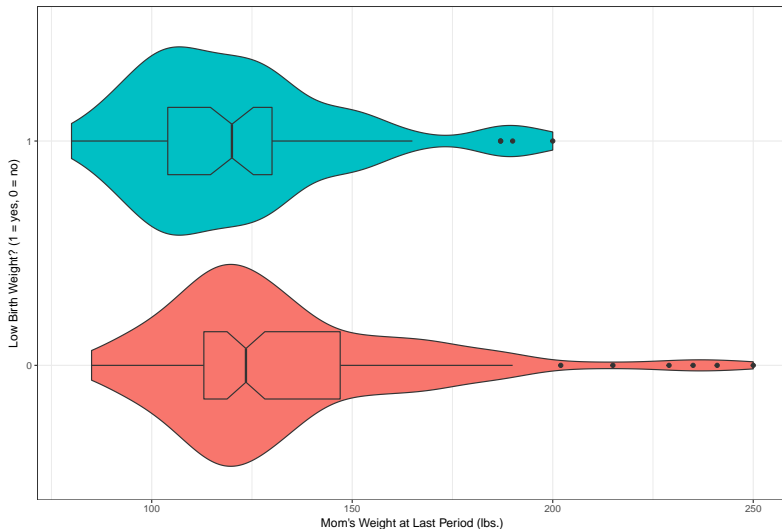
```
group variables: low
```

```
Variable type: integer
```

low	variable	missing	complete	n	mean	sd	p0	p25	median	p75	p100	hist
0	lwt	0	130	130	133.3	31.72	85	113	123.5	147	250	
1	lwt	0	59	59	122.14	26.56	80	104	120	130	200	

Can we predict $\Pr(\text{low})$ effectively with lbwt ?

Violin and Box Plots: lbw1 data



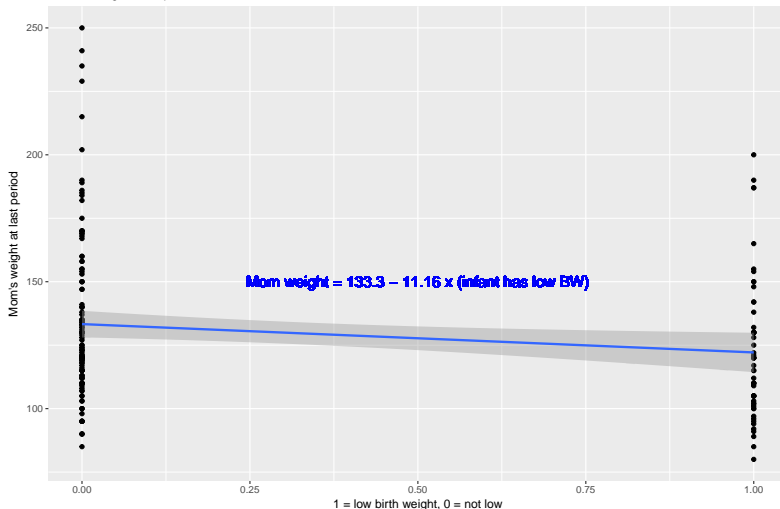
Code for Previous Slide

```
ggplot(lbw1, aes(x = factor(low), y = lwt,  
                fill = factor(low))) +  
  geom_violin() +  
  geom_boxplot(width = .3, notch = TRUE) +  
  guides(fill = FALSE) +  
  labs(x = "Low Birth Weight? (1 = yes, 0 = no)",  
       y = "Mom's Weight at Last Period (lbs.)",  
       title = "Violin and Box Plots: lbw1 data") +  
  theme_bw() +  
  coord_flip()
```


Working in Reverse: Can we predict lowt with low?

Predicting Mom's weight from low birth weight status

What is wrong with this picture?



Working in Reverse: Predicting lwt with low

Easy to go in the other direction...

```
lm(lwt ~ low, data = lbw1)
```

Call:

```
lm(formula = lwt ~ low, data = lbw1)
```

Coefficients:

(Intercept)	low
133.30	-11.16

Weight at Last Period = $133.3 - 11.16 * (\text{baby is low bw})$

- But that's reversing the outcome and predictor...

Can we fit a linear probability model? Sure, but ...

```
lm(low ~ lwt, data = lbw1)
```

Call:

```
lm(formula = low ~ lwt, data = lbw1)
```

Coefficients:

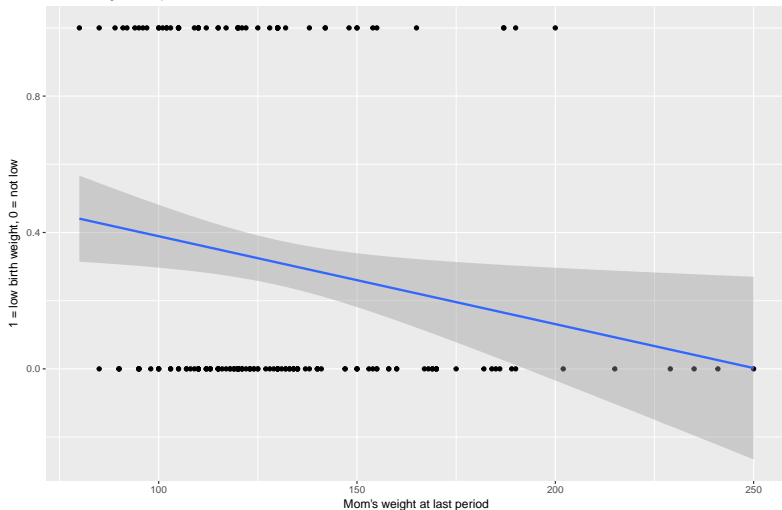
(Intercept)	lwt
0.646733	-0.002577

$\Pr(\text{low birth weight}) = 0.6467 - 0.0026 (\text{Mom's weight at last period})$

Plotting the Linear Probability Model

Linear Probability Model: $\Pr(\text{low}) = 0.6467 - 0.0026 \text{ Mom's weight}$

What is wrong with this picture?



Fitting a Model to predict a Binary Outcome

Logistic regression is the most common model used when the outcome is binary. Our response variable is assumed to take on two values - zero or one, and we then describe the probability of a “one” response, given a linear function of explanatory predictors.

- Linear regression approaches to the problem of predicting probabilities are problematic for several reasons: not least of which being that they predict probabilities greater than one and less than zero.

Logistic regression is a non-linear regression approach, since the equation for the mean of the 0/1 Y values conditioned on the values of our predictors X_1, X_2, \dots, X_k turns out to be non-linear in the β coefficients.

The Logit Link and Logistic Function

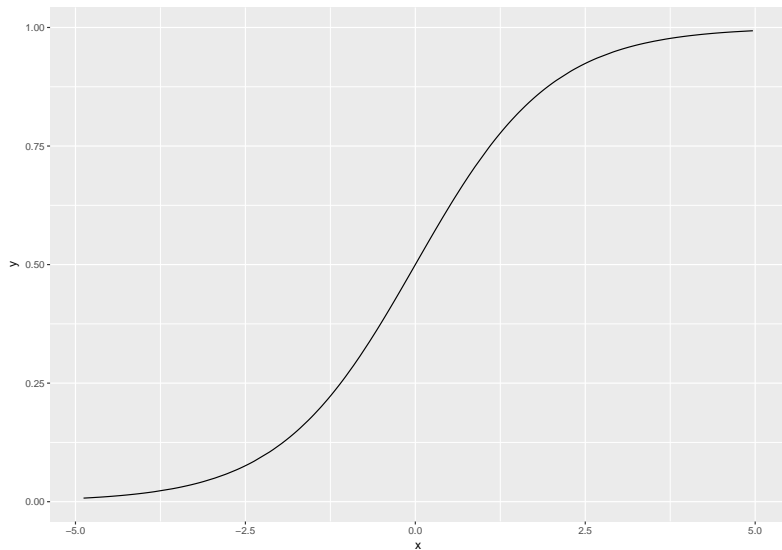
The particular link function we use in logistic regression is called the **logit link**.

$$\text{logit}(\pi) = \log\left(\frac{\pi}{1-\pi}\right) = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_k X_k$$

The inverse of the logit function is called the **logistic function**. If $\text{logit}(\pi) = \eta$, then $\pi = \frac{\exp(\eta)}{1+\exp(\eta)}$.

- The logistic function $\frac{e^x}{1+e^x}$ takes any value x in the real numbers and returns a value between 0 and 1.

The Logistic Function $y = \frac{e^x}{1+e^x}$



The logit or log odds

We usually focus on the **logit** in statistical work, which is the inverse of the logistic function.

- If we have a probability $\pi < 0.5$, then $\text{logit}(\pi) < 0$.
- If our probability $\pi > 0.5$, then $\text{logit}(\pi) > 0$.
- Finally, if $\pi = 0.5$, then $\text{logit}(\pi) = 0$.

Model 1

We'll use `glm` to get started.

```
model.1 <- glm(low ~ lwt, data = lbw1, family = binomial)
model.1
```

```
Call:  glm(formula = low ~ lwt, family = binomial, data = lbw1)
```

Coefficients:

(Intercept)	lwt
0.99831	-0.01406

Degrees of Freedom: 188 Total (i.e. Null); 187 Residual

Null Deviance: 234.7

Residual Deviance: 228.7 AIC: 232.7

Our logistic regression model

The logistic regression equation is:

$$\text{logit}(\text{Pr}(\text{low} = 1)) = \log\left(\frac{\text{Pr}(\text{low} = 1)}{1 - \text{Pr}(\text{low} = 1)}\right) = 0.99831 - 0.01406 \times \text{lwt}$$

Suppose, for instance, that we are interested in making a prediction when Mom's weight at her last period, $\text{lwt} = 130$ lbs.

So we have:

$$\text{logit}(\text{Pr}(\text{low} = 1)) = 0.99831 - 0.01406 \times 130 = -0.82949$$

Getting a Prediction from R for the Model

```
model.1 <- glm(low ~ lwt, data = lbw1, family = binomial)
```

To predict on the log odds scale, we use

```
predict(model.1, newdata = data.frame(lwt = 130))
```

```
      1  
-0.8292596
```

The Model in terms of Odds

We can exponentiate to state the odds, rather than the log odds. For a Mom at 130 lbs, we have:

$$\log \left(\frac{\text{Pr}(\text{low} = 1)}{1 - \text{Pr}(\text{low} = 1)} \right) = 0.99831 - 0.01406 \times 130 = -0.82949$$

and so we have

$$\text{Odds}(\text{low} = 1 | \text{wt} = 130) = \exp(-0.82949) = 0.4362717$$

Making a Prediction about Probability

$$\text{Odds}(\text{low} = 1 | \text{lw} = 130) = \frac{\text{Pr}(\text{low} = 1)}{1 - \text{Pr}(\text{low} = 1)} = 0.4362717$$

so

$$\text{Pr}(\text{low} = 1 | \text{lw} = 130) = \frac{\text{Odds}(\text{low} = 1 | \text{lw} = 130)}{1 + \text{Odds}(\text{low} = 1 | \text{lw} = 130)} = \frac{0.4362717}{1 + 0.4362717}$$

which is 0.304.

Obtaining a Prediction from R for Prob(low = 1)

```
model.1 <- glm(low ~ lwt, data = lbw1, family = binomial)
```

To predict on the probability scale, we can use

```
predict(model.1, newdata = data.frame(lwt = 130),  
        type = "response")
```

1

0.3038016

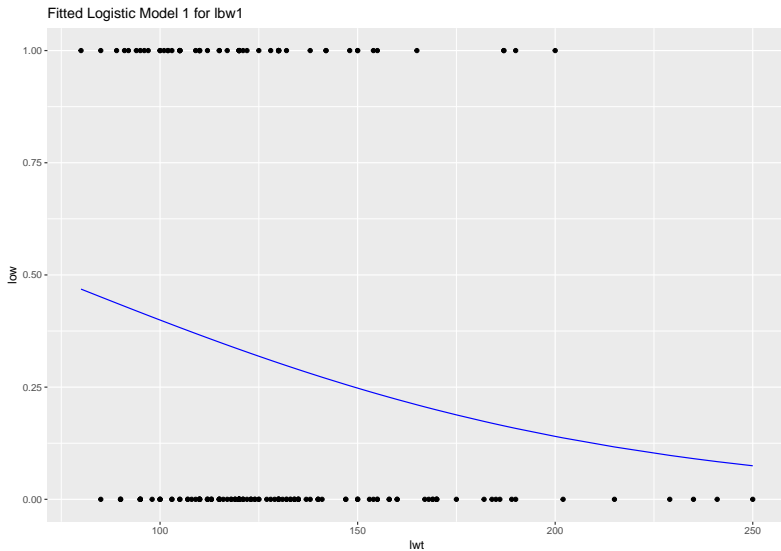
Plotting the Logistic Regression Model

We can use the `augment` function from the `broom` package to get our fitted probabilities included in the data.

```
mod1.aug <- augment(model.1, lbw1,  
                     type.predict = "response")  
  
ggplot(mod1.aug, aes(x = lwt, y = low)) +  
  geom_point() +  
  geom_line(aes(x = lwt, y = .fitted), col = "blue") +  
  labs(title = "Fitted Logistic Model 1 for lbw1")
```

- Results on next slide

Plotting the Logistic Regression Model



Cleaning up the plot

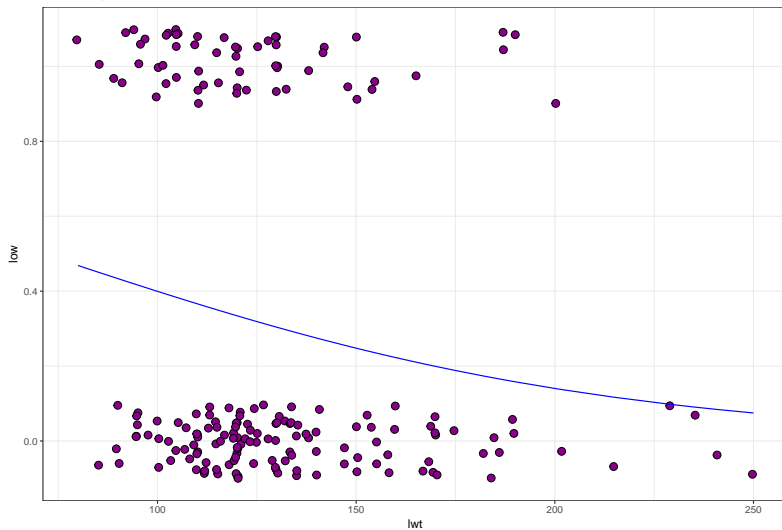
I'll add a little jitter on the vertical scale to the points, so we can avoid overlap, and also make the points a little bigger.

```
ggplot(mod1.aug, aes(x = lwt, y = low)) +  
  geom_jitter(height = 0.1, size = 3, pch = 21,  
              fill = "darkmagenta") +  
  geom_line(aes(x = lwt, y = .fitted), col = "blue") +  
  labs(title = "Fitted Logistic Model 1 for lbw1") +  
  theme_bw()
```

- Results on next slide

Cleaned up Plot of Model 1

Fitted Logistic Model 1 for lbw1



Studying the Model, Again

```
model.1
```

```
Call: glm(formula = low ~ lwt, family = binomial, data = lbw1)
```

Coefficients:

(Intercept)	lwt
0.99831	-0.01406

Degrees of Freedom: 188 Total (i.e. Null); 187 Residual

Null Deviance: 234.7

Residual Deviance: 228.7 AIC: 232.7

- $\text{logit}(\Pr(\text{low} = 1)) = 0.998 - 0.014 \text{ lwt}$
 - so ... as lwt increases, what happens to $\Pr(\text{low} = 1)$?
 - if Harry's mom weighed 130 lbs and Sally's weighed 150 lbs, how can we compare the predicted $\Pr(\text{low} = 1)$ for Harry and Sally?

Comparing Harry (lwt = 130) to Sally (lwt = 150)

```
predict(model.1, newdata = data.frame(lwt = c(130, 150)),  
       type = "response")
```

1	2
0.3038016	0.2477917

- Harry's mom weighed 130 lbs, and his predicted probability of low birth weight is 0.304
- Sally's mom weighed 150 lbs, and her predicted $\Pr(\text{low} = 1) = 0.248$

Interpreting the Coefficients of the Model

```
coef(model.1)
```

(Intercept)	lwt
0.99831432	-0.01405826

To understand the effect of lwt on low, try odds ratios.

```
exp(coef(model.1))
```

(Intercept)	lwt
2.7137035	0.9860401

Suppose Charlie's Mom weighed one pound more than Harry's.

- The **odds** of low birth weight are 0.986 times as large for Charlie as Harry.
- In general, odds ratio comparing two subjects whose lwt differ by 1 pound is 0.986

Comparing Harry to Charlie

Charlie's mom weighed 1 pound more than Harry's. The estimated odds ratio for low birth weight from the model associated with a one pound increase in 1wt is 0.986.

- If the odds ratio was 1, that would mean that Charlie and Harry had the same estimated odds of low birth weight, and thus the same estimated probability of low birth weight, despite having Moms with different weights.
- Since the odds ratio is less than 1, it means that **Charlie** has a **lower** estimated odds of low birth weight than Harry, and thus that Charlie has a lower estimated probability of low birth weight than Harry.
- If the odds ratio was greater than 1, it would mean that Charlie had a higher estimated odds of low birth weight than Harry, and thus that Charlie had a higher estimated probability of low birth weight than Harry.

The smallest possible odds ratio is ... ?

The rest of the model's output

Degrees of Freedom: 188 Total (i.e. Null); 187 Residual

Null Deviance: 234.7

Residual Deviance: 228.7 AIC: 232.7

Model	Null	Residual	Δ (model.1)
Deviance (lack of fit)	234.7	228.7	6.0
Degrees of Freedom	188	187	1

- Deviance accounted for by model.1 is 6 points on 1 df
- Can compare to a χ^2 distribution for a p value via anova

AIC = 232.7, still useful for comparing models for the same outcome

anova on a glm model

```
anova(model.1)
```

Analysis of Deviance Table

Model: binomial, link: logit

Response: low

Terms added sequentially (first to last)

	Df	Deviance	Resid. Df	Resid. Dev
NULL			188	234.67
lwt	1	5.9813	187	228.69

```
pchisq(5.9813, 1, lower.tail = FALSE)
```

```
[1] 0.01445834
```


Coming Soon

- How well does this model classify subjects?
- Receiver Operating Characteristic Curve Analysis
 - The C statistic (Area under the curve)
- Assessing Residual Plots for a Logistic Regression
- A “Kitchen Sink” Logistic Regression Model
 - Comparing Models
 - Interpreting Models with Multiple Predictors
- Logistic Regression using the `lrm` function
 - Nagelkerke R^2 , Somers' d etc.
 - Validating Summary Statistics
 - Summaries of Effects
 - Plotting In-Sample Predictions
 - Influence
 - Calibration
 - Nomograms