Generalized Linear Model & Logistic Regression

Fundamental Techniques in Data Science

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Outline

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Logistic Regression

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General Linear Model

So far, we've been discussing models with this form:

$$Y = \beta_0 + \sum_{p=1}^{P} \beta_p X_p + \varepsilon$$

This type of model is known as the *general linear model*.

- All flavors of linear regression are general linear models.
  - ANOVA
  - ANCOVA
  - Multilevel linear regression models

Components of the General Linear Model

We can break our model into pieces:
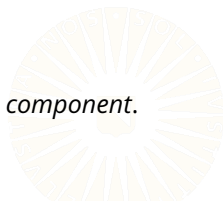
$$\eta = \beta_0 + \sum_{p=1}^{P} \beta_p X_p$$

$$Y = \eta + \varepsilon$$

Because $\varepsilon \sim N(0, \sigma^2)$, we can also write:

$$Y \sim N(\eta, \sigma^2)$$

In this representation:

- η is the *systematic component* of the model
- The normal distribution, $N(\cdot, \cdot)$, is the model's *random component*.

Components of the General Linear Model

The purpose of general linear modeling (i.e., regression modeling) is to build a model of the outcome's mean, μ_Y.

- In this case, $\mu_Y = \eta$.
- The systematic component defines the mean of Y.

The random component quantifies variability (i.e., error variance) around μ_Y.

- In the general linear model, we assume that this error variance follows a normal distribution.
- Hence the normal random component.

GENERALIZED LINEAR MODEL

Extending the General Linear Model

We can generalize the models we've been using in two important ways:

1. Allow for random components other than the normal distribution.
2. Allow for more complicated relations between μ_Y and η.
   - Allow: $g(\mu_Y) = \eta$

These extensions lead to the class of *generalized linear models* (GLMs).

Components of the Generalized Linear Model

The random component in a GLM can be any distribution from the so-called *exponential family*.

- The exponential family contains many popular distributions:
  - Normal
  - Binomial
  - Poisson
  - Many others...

The systematic component of a GLM is exactly the same as it is in general linear models:

$$\eta = \beta_0 + \sum_{p=1}^{P} \beta_p X_p$$

Link Functions

In GLMs, η does not directly describe μ_Y.

- We first transform μ_Y via a *link function*.
- $g(\mu_Y) = \eta$

The link function allows GLMs for outcomes with restricted ranges without requiring any restrictions on the range of the $\{X_p\}$.

- For strictly positive Y, we can use a *log link*:

$$\ln(\mu_y) = \eta.$$

- The general linear model employs the *identity link*:

$$\mu_y = \eta.$$

Components of the Generalized Linear Model

Every GLM is built from three components:

1. The systematic component, η.
   - A linear function of the predictors, $\{X_p\}$.
   - Describes the association between $\mathbf{X}$ and Y.

2. The link function, $g(\mu_Y)$.
   - Transforms μ_Y so that it can take any value on the real line.

3. The random component, $P(Y|g^{-1}(\eta))$
   - The distribution of the observed Y.
   - Quantifies the error variance around η.

General Linear Model as a Special Case

The general linear model is a special case of GLM.

1. Systematic component:

$$\eta = \beta_0 + \sum_{p=1}^{P} \beta_p X_p$$

2. Link function:

$$\mu_Y = \eta$$

3. Random component:

$$Y \sim \mathrm{N}(\eta, \sigma^2)$$

Example

```r
data(iris)

## General linear model:
lmFit <- lm(Petal.Length ~ Petal.Width + Species, data = iris)

## Generalized linear model:
glmFit <- glm(Petal.Length ~ Petal.Width + Species,
              family = gaussian(link = "identity"),
              data = iris)
```

Example

```
partSummary(lmFit, 2)

## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.02977 -0.22241 -0.01514  0.18180  1.17449

partSummary(glmFit, 2)

## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.02977 -0.22241 -0.01514  0.18180  1.17449
```

Example

```
partSummary(lmFit, 3)

## Coefficients:
##                    Estimate Std. Error t value Pr(>|t|)
## (Intercept)         1.21140    0.06524  18.568  < 2e-16
## Petal.Width         1.01871    0.15224   6.691 4.41e-10
## Speciesversicolor   1.69779    0.18095   9.383  < 2e-16
## Speciesvirginica    2.27669    0.28132   8.093 2.08e-13

partSummary(glmFit, 3)

## Coefficients:
##                    Estimate Std. Error t value Pr(>|t|)
## (Intercept)         1.21140    0.06524  18.568  < 2e-16
## Petal.Width         1.01871    0.15224   6.691 4.41e-10
## Speciesversicolor   1.69779    0.18095   9.383  < 2e-16
## Speciesvirginica    2.27669    0.28132   8.093 2.08e-13
```

Logistic Regression

Logistic Regression

So why do we care about the GLM when linear regression models have worked thus far?

- In a word: Classification.

In the classification task, we have a discrete, qualitative outcome.

- We will begin with the situation of two-level outcomes.
  - Alive or Dead
  - Pass or Fail
  - Pay or Default

We want to build a model that predicts class membership based on some set of interesting features.
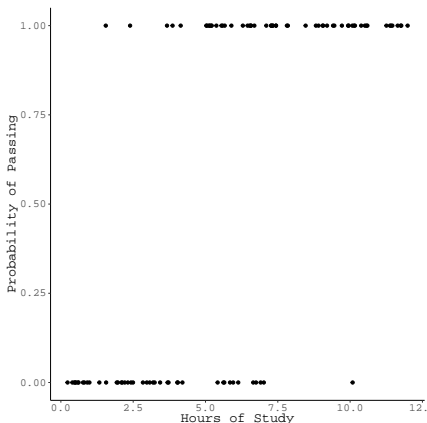
- To do so, we will use a very useful type of GLM: *logistic regression*.
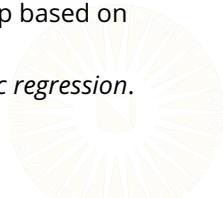
Classification Example

Suppose we want to know the effect of study time on the probability of passing an exam.
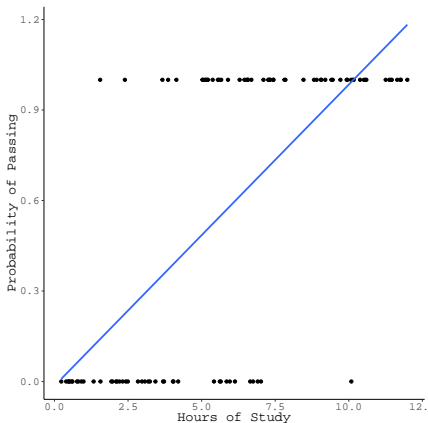
- The probability of passing must be between 0 and 1.

- We care about the probability of passing, but we only observe absolute success or failure.

  ○ $Y \in \{1, 0\}$

Linear Regression for Binary Outcomes?

What happens if we try to model these data with linear regression?

- Hmm...notice any problems?

Logistic Regression Visualized

We get a much better model using logistic regression.

- The link function ensures legal predicted values.

- The sigmoidal curve implies fluctuation in the effectiveness of extra study time.

  - More study time is most beneficial for students with around 5.5 hours of study.
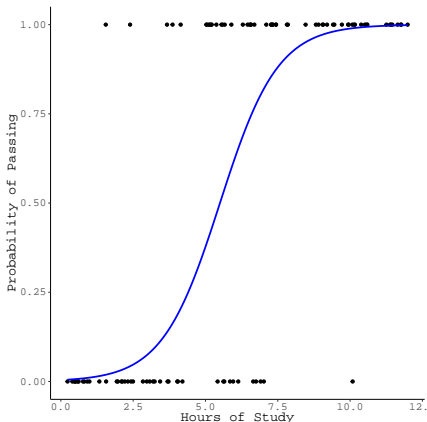
Probabilities, Odds, & Odds–Ratios

In 2017, 2535 people participated in the *Ultra-Trail du Mont-Blanc*, but only 66.55% finished the race.

- Below, you can find a cross-tabulation of finishing status and sex.

	Finish	
Sex	No	Yes
Female	95	147
Male	753	1540

- What is the *probability* of finishing for each sex?

- What are the *odds* of finishing for each sex?

- What is the *odds ratio* of finishing for males vs. females?
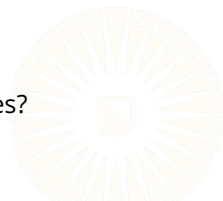
Defining the Logistic Regression Model

In logistic regression problems, we are modeling binary data:

- Usual coding: $Y \in \{1 = \text{"Success"}, 0 = \text{"Failure"}\}$.

The *Binomial* distribution is a good way to represent this kind of data.

- The systematic component in our logistic regression model will be the binomial distribution.

The mean of the binomial distribution (with $N = 1$) is the "success" probability, $\pi = P(Y = 1)$.

- We are interested in modeling $\mu_Y = \pi$:

$$g(\pi) = \beta_0 + \sum_{p=1}^{P} \beta_p X_p$$

Link Function for Logistic Regression

Because π is bounded by 0 and 1, we cannot model it directly—we must apply an appropriate link function.

- Logistic regression uses the *logit link*.

- Given π, we can define the *odds* of success as:

$$O_s = \frac{\pi}{1 - \pi}$$

- Because $\pi \in [0, 1]$, we know that $O_s \geq 0$.

- We take the natural log of the odds as the last step to fully map π to the real line.

$$\text{logit}(\pi) = \ln\left(\frac{\pi}{1 - \pi}\right)$$

Fully Specified Logistic Regression Model

Our final logistic regression model is:

$$Y \sim \text{Bin}(\pi, 1)$$

$$\text{logit}(\pi) = \beta_0 + \sum_{p=1}^{P} \beta_p X_p$$

The fitted model can be represented as:

$$\text{logit}(\hat{\pi}) = \hat{\beta}_0 + \sum_{p=1}^{P} \hat{\beta}_p X_p$$

The fitted coefficients, $\{\hat{\beta}_0, \hat{\beta}_p\}$, are interpreted in units of *log odds*.

Logistic Regression Example

If we fit a logistic regression model to the test-passing data plotted above, we get:

$$\text{logit}(\hat{\pi}_{pass}) = -3.414 + 0.683 X_{study}$$

- A student who does not study at all has -3.414 log odds of passing the exam.
- For each additional hour of study, a student's log odds of passing increase by 0.683 units.

Log odds do not lend themselves to interpretation.

- We can convert the effects back to an odds scale by exponentiation.
- $\hat{\beta}$ has log odds units, but $e^{\hat{\beta}}$ has odds units.
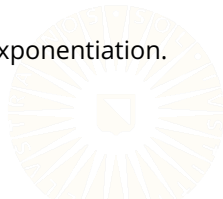
Interpretations

Exponentiating the coefficients also converts the additive effects to multiplicative effects.

- $\ln(AB) = \ln(A) + \ln(B)$
- We can interpret $\hat{\beta}$ as we would in linear regression:
  - A unit change in X_p produces an expected change of $\hat{\beta}_p$ units in $\text{logit}(\pi)$.
- After exponentiation, however, unit changes in X_p imply multiplicative changes in $O_s = \pi/(1 - \pi)$.
  - A unit change in X_p results in multiplying O_s by $e^{\hat{\beta}_p}$.

Interpretations

Exponentiating the coefficients in our toy test-passing example produces the following interpretations:

- A student who does not study is expected to pass the exam with odds of 0.033.
- For each additional hour a student studies, their odds of passing increase by 1.98 *times*.
 - Odds of passing are *multiplied* by 1.98 for each extra hour of study.

Interpretations

Exponentiating the coefficients in our toy test-passing example produces the following interpretations:

- A student who does not study is expected to pass the exam with odds of 0.033.
- For each additional hour a student studies, their odds of passing increase by 1.98 *times*.
  - Odds of passing are *multiplied* by 1.98 for each extra hour of study.

Due to the confusing interpretations of the coefficients, we often focus on the valance of the effects:

- Additional study time is associated with increased odds of passing.
- $\hat{\beta}_p > 0$ = "Increased Success", $e^{\hat{\beta}_p} > 1$ = "Increased Success"
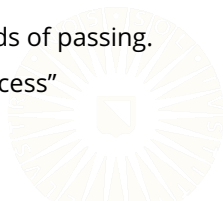
Example

Let's use logistic regression to compute the odds of finishing the UTMB.

```
## Read the UTMB data:
utmb <- readRDS(paste0(dataDir, "utmb_finish_2017.rds"))
```

We use the `glm()` function to estimate generalized linear models.

- To get a logistic regression model, we need to do two things:
  1. Specify a binary outcome variable
  2. Specify the `family = "binomial"` argument.

```
## Estimate the logistic regression model:
fit <- glm(Finish ~ Sex, family = binomial(link = "logit"), data = utmb)
```

Example

```
partSummary(fit, -1)

## Deviance Residuals:
##     Min      1Q  Median      3Q     Max
## -1.4924 -1.4924  0.8923  0.8923  0.9985
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept)   0.4366     0.1316   3.316 0.000912
## SexMale       0.2789     0.1389   2.007 0.044712
##
## (Dispersion parameter for binomial family taken to be 1)
##
##     Null deviance: 3231.3  on 2534  degrees of freedom
## Residual deviance: 3227.3  on 2533  degrees of freedom
## AIC: 3231.3
##
## Number of Fisher Scoring iterations: 4
```

Example

The raw coefficient estimates are in units of log-odds.

- We need to exponentiate the estimates to get odds ratios.

```
library(dplyr)

coef(fit) %>% exp()

## (Intercept)     SexMale
##    1.547368    1.321697
```

Multiple Logistic Regression

The preceding example was a *simple logistic regression*.

- Including multiple predictor variables in the systematic component leads to *multiple logistic regression*.
- The relative differences between simple logistic regression and multiple logistic regression are the same as those between simple linear regression and multiple linear regression.
  - The only important complication is that the regression coefficients become partial effects.

Example

Let's use logistic regression to predict the chances that Titanic passengers survived the sinking based on their age, sex, and ticket class.

```
## Read the data:
titanic <- readRDS(paste0(dataDir, "titanic.rds"))

## Estimate the logistic regression model:
fit <- glm(survived ~ age + sex + class,
           data = titanic,
           family = "binomial")
```

Example

```
partSummary(fit, -1)

## Deviance Residuals:
##     Min      1Q   Median      3Q      Max
## -2.6811  -0.6653  -0.4137   0.6367   2.4505
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept)  3.63492    0.37045   9.812  < 2e-16
## age         -0.03427    0.00716  -4.787 1.69e-06
## sexmale     -2.58872    0.18701 -13.843  < 2e-16
## class2nd    -1.19911    0.26158  -4.584 4.56e-06
## class3rd    -2.45544    0.25322  -9.697  < 2e-16
##
## (Dispersion parameter for binomial family taken to be 1)
##
##     Null deviance: 1182.77  on 886  degrees of freedom
## Residual deviance: 801.59  on 882  degrees of freedom
## AIC: 811.59
##
## Number of Fisher Scoring iterations: 5
```

Example

Compute odds ratios.

```
(or <- coef(fit) %>% exp())

## (Intercept)       age     sexmale    class2nd    class3rd
## 37.8988400  0.9663058  0.0751161  0.3014609  0.0858252
```

Odds ratios smaller than 1.0 can be difficult to explain.

- We can ease interpretation by reciprocating the estimates.

```
1 / or

## (Intercept)       age     sexmale    class2nd    class3rd
## 0.02638603  1.03486914 13.31272574  3.31717996 11.65158920
```

Example

To convince ourselves that the above operation is sensible, we can compare the inverse odds ratios to the odds ratios we get from predicting the chances of dying.

```
library(magrittr)

fit2 <- titanic %>%
    mutate(died = relevel(survived, ref = "yes")) %$%
    glm(died ~ age + sex + class, family = "binomial")
```

Example

```
partSummary(fit2, -1)

## Deviance Residuals:
##     Min      1Q   Median      3Q      Max
## -2.4505  -0.6367   0.4137   0.6653   2.6811
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -3.63492    0.37045  -9.812  < 2e-16
## age          0.03427    0.00716   4.787 1.69e-06
## sexmale      2.58872    0.18701  13.843  < 2e-16
## class2nd     1.19911    0.26158   4.584 4.56e-06
## class3rd     2.45544    0.25322   9.697  < 2e-16
##
## (Dispersion parameter for binomial family taken to be 1)
##
##     Null deviance: 1182.77  on 886  degrees of freedom
## Residual deviance:  801.59  on 882  degrees of freedom
## AIC: 811.59
##
## Number of Fisher Scoring iterations: 5
```

Example

We get the same odds ratios that we derived through reciprocation.

```
coef(fit2) %>% exp()

## (Intercept)          age      sexmale     class2nd     class3rd
##  0.02638603   1.03486914  13.31272574   3.31717996  11.65158920

1 / or

## (Intercept)          age      sexmale     class2nd     class3rd
##  0.02638603   1.03486914  13.31272574   3.31717996  11.65158920
```

Example in Equations

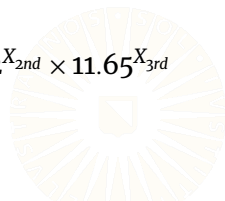Here's the symbolic representation of our logistic regression model:

$$\text{logit}(\pi_{died}) = \beta_0 + \beta_1 X_{age} + \beta_2 X_{male} + \beta_3 X_{2nd} + \beta_4 X_{3rd}$$

By fitting this model to the *titanic* data we get:

$$\text{logit}(\hat{\pi}_{died}) = -3.63 + 0.03 X_{age} + 2.59 X_{male} + 1.2 X_{2nd} + 2.46 X_{3rd}$$

Exponentiating the coefficients produces:

$$\frac{\hat{\pi}_{died}}{1 - \hat{\pi}_{died}} = \frac{\hat{\pi}_{died}}{\hat{\pi}_{survived}} = 0.03 \times 1.03^{X_{age}} \times 13.31^{X_{male}} \times 3.32^{X_{2nd}} \times 11.65^{X_{3rd}}$$

Exponentiating the Systematic Component

$$\text{logit}(\hat{\pi}_{died}) = -3.63 + 0.03X_{age} + 2.59X_{male} + 1.2X_{2nd} + 2.46X_{3rd}$$

$$e^{\text{logit}(\hat{\pi}_{died})} = e^{\left(-3.63 + 0.03X_{age} + 2.59X_{male} + 1.2X_{2nd} + 2.46X_{3rd}\right)}$$

$$\frac{\hat{\pi}_{died}}{\hat{\pi}_{survived}} = e^{-3.63} \times e^{0.03X_{age}} \times e^{2.59X_{male}} \times e^{1.2X_{2nd}} \times e^{2.46X_{3rd}}$$

$$= e^{-3.63} \times \left(e^{0.03}\right)^{X_{age}} \times \left(e^{2.59}\right)^{X_{male}} \times \left(e^{1.2}\right)^{X_{2nd}} \times \left(e^{2.46}\right)^{X_{3rd}}$$

$$= 0.03 \times 1.03^{X_{age}} \times 13.31^{X_{male}} \times 3.32^{X_{2nd}} \times 11.65^{X_{3rd}}$$

CLASSIfiCATION

Predictions from Logistic Regression

Given a fitted logistic regression model, we can get predictions for new observations of $\{X_p\}$, $\{X'_p\}$.

- Directly applying $\{\hat{\beta}_0, \hat{\beta}_p\}$ to $\{X'_p\}$ will produce predictions on the scale of η:

$$\hat{\eta}' = \hat{\beta}_0 + \sum_{p=1}^{P} \hat{\beta}_p X'_p$$

- By applying the inverse link function, $g^{-1}(\cdot)$, to $\hat{\eta}'$, we get predicted success probabilities:

$$\hat{\pi}' = g^{-1}(\hat{\eta}')$$

Predictions from Logistic Regression

In logistic regression, the inverse link function, $g^{-1}(\cdot)$, is the *logistic function*:

$$\text{logistic}(X) = \frac{e^X}{1 + e^X}$$

So, we convert $\hat{\eta}'$ to $\hat{\pi}'$ by:

$$\hat{\pi}' = \frac{e^{\hat{\eta}'}}{1 + e^{\hat{\eta}'}} = \frac{\exp\left(\hat{\beta}_0 + \sum_{p=1}^{P} \hat{\beta}_p X'_p\right)}{1 + \exp\left(\hat{\beta}_0 + \sum_{p=1}^{P} \hat{\beta}_p X'_p\right)}$$

Classification with Logistic Regression

Once we have computed the predicted success probabilities, $\hat{\pi}'$, we can use them to classify new observations.

- By choosing a threshold on $\hat{\pi}'$, say $\hat{\pi}' = t$, we can classify the new observations as "Successes" or "Failures":

$$\hat{Y}' = \left\{ \begin{array}{ll} 1 & if \quad \hat{\pi}' \geq t \\ 0 & if \quad \hat{\pi}' < t \end{array} \right.$$
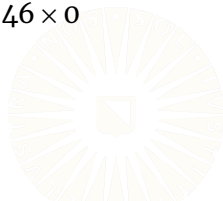
Classification Example

Say we want to classify a hypothetical passenger as either having died or survived the sinking.

- Assume this passenger has the following characteristics:
  - They are 17 years old
  - They are male
  - They are a second class passenger

First we plug their predictor data into the fitted model to get their model-implied η:

$$\hat{\eta}_{died} = -3.63 + 0.03 \times 17 + 2.59 \times 1 + 1.2 \times 1 + 2.46 \times 0$$

$$= 0.736$$

Classification Example

Next we convert the predicted η value into a model-implied success probability by applying the logistic function:

$$\frac{e^{0.736}}{1 + e^{0.736}} = 0.676$$

Finally, to make the classification, assume a threshold of $\hat{\pi}' = 0.5$ as the decision boundary.

- Because $0.676 > 0.5$ we would classify this passenger as having died.

Example

Let's generate a grid of hypothetical passengers to explore the model's predictions.

```
passengers <-
    expand.grid(
        age   = 1:100,
        sex   = c("male", "female"),
        class = c("1st", "2nd", "3rd")
    ) %>%
    data.frame()
```

Example

View 10 random passengers.

```
passengers %>% slice_sample(n = 10)

##    age    sex class
## 1   68   male   1st
## 2   51   male   3rd
## 3   62 female   2nd
## 4   40 female   2nd
## 5   60 female   2nd
## 6   50   male   2nd
## 7   63   male   1st
## 8   54   male   3rd
## 9   38   male   3rd
## 10  48   male   3rd
```

Example

We can generate predictions on the scale of the linear predictor (i.e., log-odds), or we can generate predicted probabilities.

```
passengers %<>%
    mutate(
        ## Predicted log odds of dying:
        etaHat = predict(fit2, newdata = ., type = "link"),

        ## Predicted probabilities of dying:
        piHat = predict(fit2, newdata = ., type = "response")
    )
```

We can then use the predicted probabilities of dying to classify:

```
passengers %<>%
    mutate(dieHat = ifelse(piHat > 0.5, "dead", "alive") %>% factor())
```

Example

View the predictions for 10 random passengers:

```
passengers %>% slice_sample(n = 10)

##     age    sex class      etaHat     piHat  dieHat
## 1    21   male   1st -0.32642546 0.4191106   alive
## 2     1   male   2nd  0.18718989 0.5466613    dead
## 3    60   male   2nd  2.20941388 0.9010917    dead
## 4    12 female   2nd -2.02450570 0.1166539   alive
## 5     7   male   3rd  1.64916736 0.8387785    dead
## 6    92   male   1st  2.10709832 0.8915912    dead
## 7    75   male   3rd  3.97986619 0.9816547    dead
## 8    28 female   2nd -1.47610597 0.1860163   alive
## 9    52 female   3rd  0.60282119 0.6463015    dead
## 10   72 female   2nd  0.03199328 0.5079976    dead
```
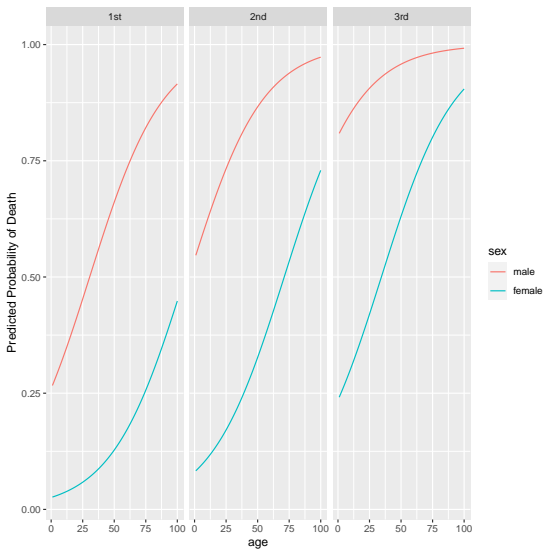
Example

We can visualize the predicted probabilities of dying.

```
library(ggplot2)

ggplot(passengers, aes(age, piHat, color = sex)) +
    geom_line() +
    facet_wrap(vars(class)) +
    ylab("Predicted Probability of Death")
```

Example

Example

We can also visualize the classifications the model would make.

```
ggplot(passengers, aes(dieHat, age, color = class)) +
    geom_boxplot() +
    facet_wrap(vars(sex)) +
    xlab("Predicted Death")
```

Example