```
## Logistic regression
This far we have used the `lm' function to fit our regression models.
    `lm' is great, but limitedâin particular it only fits models for
    continuous dependent variables. For categorical dependent variables we
##
##
    can use the `glm()' function.
##
    For these models we will use a different dataset, drawn from the
    National Health Interview Survey. From the [CDC website]:
##
##
          The National Health Interview Survey (NHIS) has monitored
##
          the health of the nation since 1957. NHIS data on a broad
          range of health topics are collected through personal
##
          household interviews. For over 50 years, the U.S. Census
          Bureau has been the data collection agent for the National
##
##
          Health Interview Survey. Survey results have been
##
          instrumental in providing data to track health status,
##
          health care access, and progress toward achieving national
##
          health objectives.
    Load the National Health Interview Survey data:
NH11 <- readRDS("dataSets/NatHealth2011.rds")</pre>
labs <- attributes(NH11) $labels
    [CDC website] http://www.cdc.gov/nchs/nhis.htm
## Logistic regression example
Let's predict the probability of being diagnosed with hypertension
    based on age, sex, sleep, and bmi
str(NH11$hypev) # check stucture of hypev
levels(NH11$hypev) # check levels of hypev
# collapse all missing values to NA
NH11$hypev <- factor(NH11$hypev, levels=c("2 No", "1 Yes"))</pre>
# run our regression model
\label{limits} \verb|hyp.out <- glm(hypev~age_p+sex+sleep+bmi,
             data=NH11, family="binomial")
coef(summary(hyp.out))
## Logistic regression coefficients
Generalized linear models use link functions, so raw coefficients are
##
    difficult to interpret. For example, the age coefficient of .06 in the
    previous model tells us that for every one unit increase in age, the
##
    log odds of hypertension diagnosis increases by 0.06. Since most of us
##
    are not used to thinking in log odds this is not too helpful!
    One solution is to transform the coefficients to make them easier to
    interpret
hyp.out.tab <- coef(summary(hyp.out))</pre>
hyp.out.tab[, "Estimate"] <- exp(coef(hyp.out))</pre>
hyp.out.tab
## Generating predicted values
In addition to transforming the log-odds produced by `glm' to odds, we
##
    can use the `predict()' function to make direct statements about the
##
    predictors in our model. For example, we can ask "How much more likely
##
     is a 63 year old female to have hypertension compared to a 33 year old
    female?".
# Create a dataset with predictors set at desired levels
predDat <- with (NH11.
               expand.grid(age_p = c(33, 63),
                               = "2 Female",
                           sex :
                           bmi = mean(bmi, na.rm = TRUE),
                           sleep = mean(sleep, na.rm = TRUE)))
# predict hypertension at those levels
cbind(predDat, predict(hyp.out, type = "response",
                      se.fit = TRUE, interval="confidence",
                      newdata = predDat))
    This tells us that a 33 year old female has a 13% probability of
##
    having been diagnosed with hypertension, while and 63 year old female
##
    has a 48% probability of having been diagnosed.
```

Regression with binary outcomes

Packages for $% \left(1\right) =\left(1\right) ^{2}$ computing and graphing predicted values

##

the model.