

Unit2

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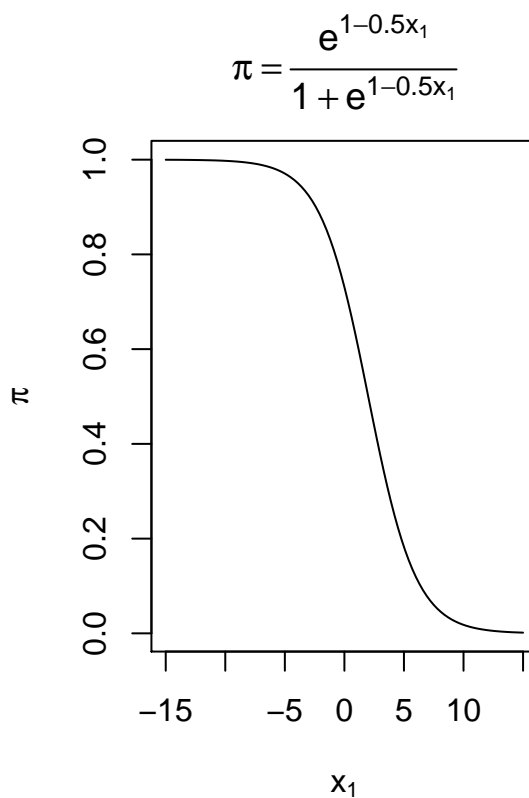
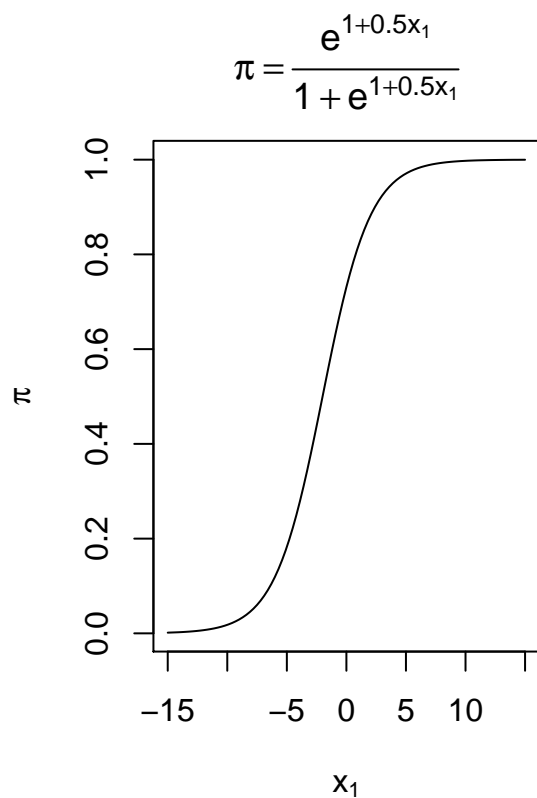
September 13, 2017

Plot the logistic regression model

```
par(mfrow = c(1, 2))
beta0 = 1
beta1 = 0.5

curve(expr = exp(beta0 + beta1*x) / (1 + exp(beta0 + beta1*x)),
      xlim = c(-15, 15), col = "black",
      main = expression(pi == frac(e^{1 + 0.5*x[1]}, 1+e^{1+0.5*x[1]})),
      xlab = expression(x[1]), ylab = expression(pi))

beta1 = -0.5
curve(expr = exp(beta0 + beta1*x) / (1 + exp(beta0 + beta1*x)),
      xlim = c(-15, 15), col = "black",
      main = expression(pi == frac(e^{1 - 0.5*x[1]}, 1+e^{1 - 0.5*x[1]})),
      xlab = expression(x[1]), ylab = expression(pi))
```



Example

```
df = read.table(file = "Placekick.csv", header = TRUE, sep = ",")
str(df)
```

```
## 'data.frame': 1425 obs. of 9 variables:
## $ week : int 1 1 1 1 1 1 1 1 1 1 ...
## $ distance: int 21 21 20 28 20 25 20 27 44 32 ...
## $ change : int 1 0 0 0 0 0 0 1 1 0 ...
## $ elap30 : num 24.72 15.85 0.45 13.55 21.87 ...
## $ PAT : int 0 0 1 0 1 0 1 0 0 0 ...
## $ type : int 1 1 1 1 0 0 0 0 0 0 ...
## $ field : int 1 1 1 1 0 0 0 0 0 0 ...
## $ wind : int 0 0 0 0 0 0 0 0 0 0 ...
## $ good : int 1 1 1 1 1 1 1 1 1 1 ...
```

```
head(df)
```

```
##   week distance change elap30 PAT type field wind good
## 1    1      21      1 24.7167  0   1    1    0    1
## 2    1      21      0 15.8500  0   1    1    0    1
## 3    1      20      0  0.4500  1   1    1    0    1
## 4    1      28      0 13.5500  0   1    1    0    1
## 5    1      20      0 21.8667  1   0    0    0    1
## 6    1      25      0 17.6833  0   0    0    0    1
```

```
# check NA in each column
```

```
apply(is.na(df), 2, sum)
```

```
##   week distance change elap30 PAT type field wind
##    0         0      0      0    0    0    0    0
##   good
##    0
```

```
# check the dependent variable of interest
```

```
table(df$good)
```

```
##
##    0    1
## 163 1262
```

```
prop.table(table(df$good))
```

```
##
##      0      1
## 0.114386 0.885614
```

use distance explanatory variable to estimate the probability of a successful placekick

```
mod.fit = glm(formula = good ~ distance, family = binomial(link = logit), data = df)
mod.fit
```

```
##
## Call:  glm(formula = good ~ distance, family = binomial(link = logit),
```

```
##      data = df)
##
## Coefficients:
## (Intercept)      distance
##      5.812      -0.115
##
## Degrees of Freedom: 1424 Total (i.e. Null);  1423 Residual
## Null Deviance:      1013
## Residual Deviance: 775.7      AIC: 779.7

logit( $\pi$ ) = 5.812 - 0.115distance

# there are many information stored within the mod.fit object
names(mod.fit)

## [1] "coefficients"      "residuals"          "fitted.values"
## [4] "effects"           "R"                   "rank"
## [7] "qr"                 "family"              "linear.predictors"
## [10] "deviance"           "aic"                 "null.deviance"
## [13] "iter"               "weights"              "prior.weights"
## [16] "df.residual"         "df.null"              "y"
## [19] "converged"          "boundary"             "model"
## [22] "call"               "formula"              "terms"
## [25] "data"               "offset"               "control"
## [28] "method"             "contrasts"            "xlevels"

length(mod.fit$coefficients)

## [1] 2

mod.fit$coefficients

## (Intercept)      distance
##      5.8120798    -0.1150267

mod.fit$coefficients[1]

## (Intercept)
##      5.81208

summary(object = mod.fit)

##
## Call:
## glm(formula = good ~ distance, family = binomial(link = logit),
##      data = df)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7441    0.2425    0.2425    0.3801    1.6092
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  5.812080   0.326277   17.81  <2e-16 ***
## distance     -0.115027   0.008339  -13.79  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```

## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 1013.43 on 1424 degrees of freedom
## Residual deviance: 775.75 on 1423 degrees of freedom
## AIC: 779.75
##
## Number of Fisher Scoring iterations: 6

class(mod.fit)

## [1] "glm" "lm"

methods(class = glm)

## [1] add1          anova          coerce         confint
## [5] cooks.distance deviance       drop1          effects
## [9] extractAIC     family        formula       influence
## [13] initialize     logLik        model.frame   nobs
## [17] predict        print         residuals     rstandard
## [21] rstudent       show          slotsFromS3   summary
## [25] vcov           weights
## see '?methods' for accessing help and source code

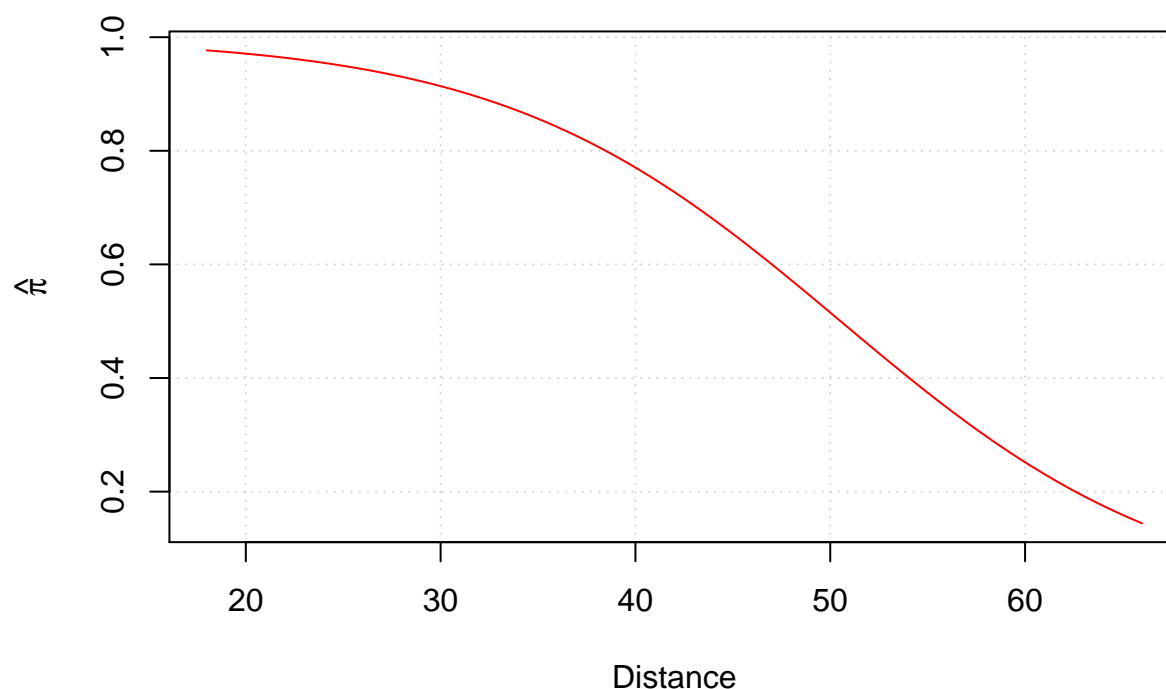
summary(df$distance)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 18.00 20.00 20.00 27.55 36.00 66.00

curve(expr = exp(mod.fit$coefficients[1] + mod.fit$coefficients[2]*x) /
      (1 + exp(mod.fit$coefficients[1] + mod.fit$coefficients[2]*x)),
      col = "red", xlim = c(18, 66), ylab = expression(hat(pi)), xlab = "Distance",
      main = "Estimated probability of success for a placekick", panel.first = grid()
      )

```

Estimated probability of success for a placekick



```
mod.fit2 = glm(formula = good ~ change + distance, family = binomial(link = logit), data = df)
summary(mod.fit2)
```

```
##
## Call:
## glm(formula = good ~ change + distance, family = binomial(link = logit),
##      data = df)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7061   0.2282   0.2282   0.3750   1.5649
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  5.893181   0.333184  17.687  <2e-16 ***
## change       -0.447783   0.193673  -2.312   0.0208 *
## distance     -0.112889   0.008444 -13.370  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 1013.4  on 1424  degrees of freedom
## Residual deviance:  770.5  on 1422  degrees of freedom
## AIC: 776.5
##
```

```

## Number of Fisher Scoring iterations: 6
mod.fit2$coefficients

## (Intercept)      change      distance
##   5.8931814   -0.4477832   -0.1128888

It takes 6 iterations to come up with the parameters

newdata = data.frame(change = 0.5, distance = 50)
predict(mod.fit2, newdata, type="response")

##           1
## 0.5062124

b0 = mod.fit2$coefficients[1]
b1 = mod.fit2$coefficients[2]
b2 = mod.fit2$coefficients[3]

summary(df$change)

##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.
## 0.0000 0.0000 0.0000 0.2519 1.0000 1.0000

x_range = seq(from = min(df$distance), to=max(df$distance), by = .1)

curve1 = exp(b0 + b1*0.5 + b2*x_range) /
  (1 + exp(b0 + b1*0.5 + b2*x_range))

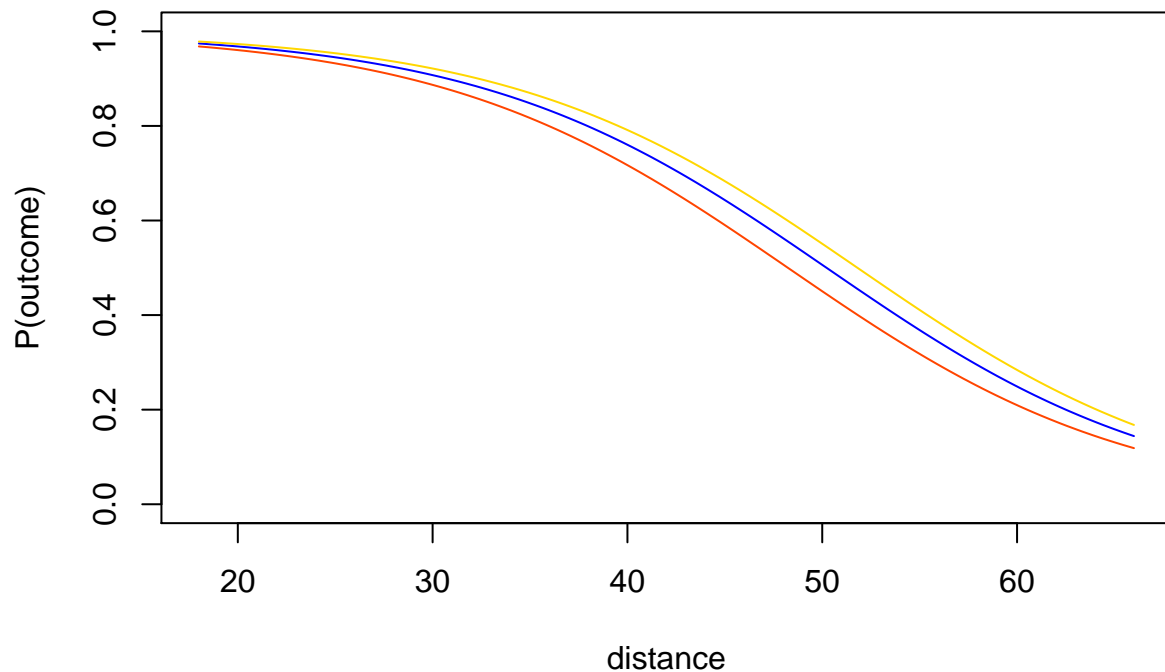
curve2 = exp(b0 + b1*0.1 + b2*x_range) /
  (1 + exp(b0 + b1*0.1 + b2*x_range))

curve3 = exp(b0 + b1*1 + b2*x_range) /
  (1 + exp(b0 + b1*1 + b2*x_range))

plot(x_range, curve1, ylim = c(0, 1), col = "blue", type= "l", xlab = "distance",
      ylab = "P(outcome)", main = "Probability of Success for a Placekick")
lines(x_range, curve2, col = "gold", type= "l")
lines(x_range, curve3, col = "orangered", type= "l")

```

Probability of Success for a Placekick



The blue line is with change = 0.5, the red line 1.0 and the yellow line 0.1. The line with smaller change value has higher probability of success.

Variance - Covariance Matrix

```
vcov(mod.fit2)
```

```
##           (Intercept)      change      distance
## (Intercept) 0.111011379 -0.0094878323 -2.625598e-03
## change      -0.009487832  0.0375091687 -1.311512e-04
## distance     -0.002625598 -0.0001311512  7.129494e-05
```

```
summary(mod.fit2)
```

```
##
## Call:
## glm(formula = good ~ change + distance, family = binomial(link = logit),
##      data = df)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -2.7061   0.2282   0.2282   0.3750   1.5649
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  5.893181   0.333184  17.687  <2e-16 ***
```

```
## change      -0.447783    0.193673  -2.312    0.0208 *
## distance    -0.112889    0.008444 -13.370    <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 1013.4  on 1424  degrees of freedom
## Residual deviance:  770.5  on 1422  degrees of freedom
## AIC: 776.5
##
## Number of Fisher Scoring iterations: 6
```

```
library(car)
Anova(mod = mod.fit2, test = "LR")
```

```
## Analysis of Deviance Table (Type II tests)
##
## Response: good
##      LR Chisq Df Pr(>Chisq)
## change      5.246  1      0.022 *
## distance 218.650  1      <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

when `Anova()` is not available with more complex models than logistic regression

```
mod.fit2 = glm(formula = good ~ change + distance, family = binomial(link = logit), data = df)
mod.fit.Ho = glm(formula = good ~ distance, family = binomial(link = logit), data = df)
anova(mod.fit.Ho, mod.fit2, test = "Chisq") # note "a" nova, not Anova
```

```
## Analysis of Deviance Table
##
## Model 1: good ~ distance
## Model 2: good ~ change + distance
##   Resid. Df Resid. Dev Df Deviance Pr(>Chi)
## 1      1423      775.75
## 2      1422      770.50  1    5.2455    0.022 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

General approach for log-likelihood function

check page 71 textbook

```
logL = function(beta, x, Y) {
  pi = exp(beta[1] + beta[2] * x) / ( 1 + exp(beta[1] + beta[2] * x))
  sum(Y*log(pi) + (1-Y)*log(1-pi))
}

logL(beta = mod.fit$coefficients, x = df$distance, Y = df$good)
```

```
## [1] -387.8725
# find starting values for parameter estimate
reg.mod = lm(formula = good ~ distance, data = df)
```



```

reg.mod$coefficients

## (Intercept)    distance
##  1.25202444 -0.01330212

# use control = list(fnscale = -1) to "maximize" instead of minimize
mod.fit.optim = optim(par = reg.mod$coefficients, fn = logL,
                      hessian = TRUE, x = df$distance, Y = df$good, control = list(fnscale = -1),
                      method = "BFGS")
names(mod.fit.optim)

## [1] "par"          "value"          "counts"         "convergence" "message"
## [6] "hessian"

mod.fit.optim$par

## (Intercept)    distance
##   5.8112544   -0.1150046

mod.fit.optim$convergence

## [1] 0

-solve(mod.fit.optim$hessian)

##              (Intercept)      distance
## (Intercept)  0.106482867 -2.607258e-03
## distance    -0.002607258  6.957463e-05

```

Deviance

```

mod.fit2 = glm(formula = good ~ change + distance, family = binomial(link = logit), data = df)
mod.fit.Ho = glm(formula = good ~ distance, family = binomial(link = logit), data = df)

dframe = mod.fit.Ho$df.residual - mod.fit2$df.residual

stat = mod.fit.Ho$deviance - mod.fit2$deviance

pvalue = 1 - pchisq(q= stat, df = dframe)
data.frame(Ho.resid.dev = mod.fit2$deviance, Ha.resid.dev = mod.fit2$deviance,
            df = dframe, stat = round(stat, 4),
            pvalue = round(pvalue, 4))

##   Ho.resid.dev Ha.resid.dev df   stat pvalue
## 1      770.4995      770.4995 1 5.2455 0.022

```