1.3.2 Additive Errors Nonlinear Model

Previous example had O linear trend, @ Non-Gaussian enors.

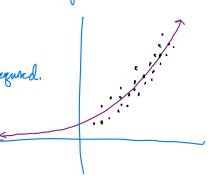
Example: exposed of

Nonlinear additive model:

$$Y_i = g(X_i, \underline{B}) + \varepsilon_i$$

Usually $\mathcal{E}_i \sim N(0, 6^2)$ but $g(X_i, \beta) \neq x_i^T \beta \Rightarrow \text{more, likelihood reguned.}$

1) non-linear trend , 2) Gaussian errors.



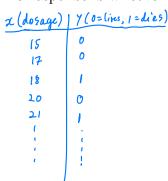
1.3.3 Generalized Linear Models

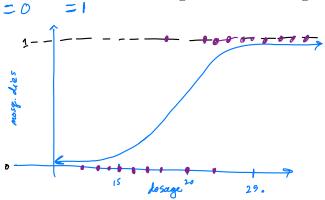
Regression: build a relationship w/ parameter (wear) + covarietes.

LM: stochastic elevent is additive v/mean.

GLM: Stochastic elevat be different.

Imagine an experiment where individual mosquitos are given some dosage of pesticide. The response is whether the mosquito lives or dies. The data might look something like:





Goal: Model the relationship between the predictor and response.

sounds like regression!

dosage dead/alive

Big differee: Y's are not continuous. They only take valves of O or 1 (binary response).

Question: What would a curve of best fit look like? Would it only take values of 0 or 1?

It seems sensible to have a curve which takes values near O for low doses, near I for high doses and intermediate values between O and I for intermediate doses.

What does this curve represent? Probability.

Refined Goal: Model teletionship between predictor (do sage) I probability (of mosq. dies).

Let's build a sensible model. Note: We do not observe the probability!

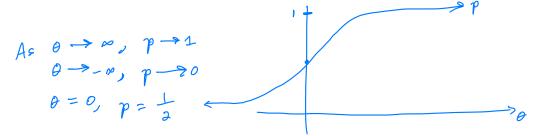
If I flip a possibly bised air of 3 heads do you know P(heads) = .75?

Step 1: Find a function that behaves the way we want.

Ly like the blue curre.

Consider the Logistic function

$$p = \frac{\exp(\theta)}{1 + \exp(\theta)}$$



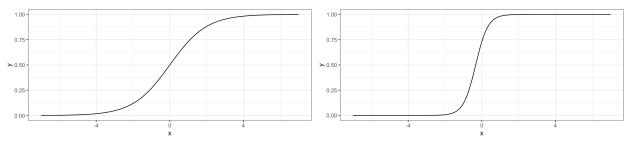


By changing &, we can change location, slope, direction of this function.

$$\partial = \beta_0 + \beta_1 \times \Rightarrow p = \frac{\exp(\beta_0 + \beta_1 \times)}{1 + \exp(\beta_0 + \beta_1 \times)}$$

```
# understanding the logistic function
# first, theta just equals x
x <- seq(-7, 7, .1)
theta <- x
y <- exp(theta)/(1 + exp(theta))
ggplot() + geom_line(aes(x, y))

# now, let theta be a linear function of x
theta <- 1 + 3*x
y <- exp(theta)/(1 + exp(theta))
ggplot() + geom_line(aes(x, y))</pre>
```



Now we have the ability to connect probabilities to covariate ac! We'd be done if we observed probabilities, but our response only takes values of D or 1.

Step 2: Build a stochastic mechanism to relate to a binary response.

Recall the Benoulli distribution:

$$y = \begin{cases} 0 & v.p. & 1-p \\ 1 & v.p. & p \end{cases}$$

Dade to biased win flip example w/ p=0.75. Flip once, You will get O(tails) or 1 (heads).

Aside: You may be more familiar L/ Binomial distribution, which counts # of successes for notices X takes values in $\{0,1,\ldots,n\}$ $X = \sum_{i=1}^{n} Y_i, \ Y_i \stackrel{iid}{\sim} Bemouth (p).$ $P(X=k) = \binom{n}{k} p^k (1-p)^{n+k}$

Step 3: Put Step 1 and Step 2 together.

V. N Bemouilli(
$$p_i$$
)

N prob. of ith observed)

OR

(observed)

Pi = $\frac{\exp(\theta_i)}{1 + \exp(\theta_i)}$, $\theta_i = \beta_0 + \beta_i x_i$

(unobserved)

Pi = $\frac{\exp(\beta_0 + \beta_i x_i)}{1 + \exp(\beta_0 + \beta_i x_i)}$

Goal: estimate Bo + B1. Find the "best" estimates.

Fitting our model: Does OLS make sense? A)O.

What else can we do? Maximum likelihood!

L> Find the parameters (p's) which make the density agree best a/ data!

Consider the likelihood contribution.

$$L_i(p_i|Y_i) = p_i^{\gamma_i} \left(1-p_i\right)^{\lfloor \gamma_i \rfloor} \quad \left(\gamma_i^{\prime}\right)^{s} \quad \text{are on 1}.$$

So the log-likelihood contribution is

Manipulating:
$$p_{i} + p_{i} \exp(\theta_{i}) = \exp(\theta_{i})$$

$$p_{i} = (1-p_{i}) \exp(\theta_{i})$$

$$p_{i} \exp(-\theta_{i}) = 1-p_{i}$$

$$\exp(\theta_{i}) = \exp(\theta_{i}) = 1-p_{i}$$

$$\exp(\theta_{i}) = 1-p_{i}$$

Plugging (1) + (2) into (2) Which gives us,

$$\ell_i(\theta_i) = -\log(1+\exp(0i)) + \gamma_i \theta_i \quad \text{(now in terms of θ_i not p_i)}.$$

$$\text{Not a coincident!} \quad \text{Because the "sensible function" $p_i = \frac{\exp(0i)}{1+\exp(0i)} \text{ works well together}.}$$

So the log-likelihood is

$$\ell(\theta_1, \dots, \theta_n) = \sum_{i=1}^n \text{li}(\theta_i)$$

$$= \sum_{i=1}^n \left\{ -\log\left(1 + \exp(\theta_i)\right) + \gamma_i \theta_i \right\}$$

$$\Rightarrow l(\beta_0,\beta_1) = \sum_{i=1}^{n} \left\{ -log(1+exp(\beta_0+\beta_ix_i)) + y_i(\beta_0+\beta_ix_i) \right\}$$

To optimize? Must be hore numerically.

```
## data on credit default
         data("Default", package = "ISLR")
         head(Default)
##
     default student
                         balance
                                     income
## 1
          No
                   No 729.5265 44361.625
## 2
          No
                  Yes 817.1804 12106.135
## 3
                   No 1073.5492 31767.139
          No
## 4
          No
                   No 529.2506 35704.494
## 5
                   No 785.6559 38463.496
          No
## 6
          No
                  Yes 919.5885 7491.559
         ## fit model with ML
         m0 <- glm(default ~ balance, data = Default, family =
                 Toptimizing likelihood numerally.
binomial)
         tidy(m0) [> kable()
                          Bome Bi, mis sol (B.), sol(B.)
                            estimate std.error
             term
                                                 statistic p.value
             (Intercept) -10.6513306 0.3611574 -29.49221
                                                             0
                         0.0054989 \ 0.0002204 \ 24.95309
                                                             0
             balance
         glance(m0) |> kable()
 null.deviance
               df.null
                                    AIC
                                              BIC deviance
                                                             df.residual
                         logLik
                                                                         nobs
                9999 -798.2258 1600.452 1614.872 1596.452
     2920.65
                                                                  9998 10000
```

```
## plot the curve

x_new <- seq(0, 2800, length.out = 200)

theta <- m0$coefficients[1] + m0$coefficients[2]*x_new

p_hat <- exp(theta)/(1 + exp(theta))

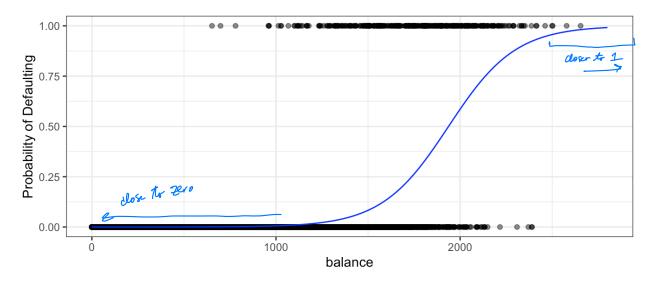
(strind possibitive ggplot() +

geom_point(aes(balance, as.numeric(default) - 1), alpha =

0.5, data = Default) +

Change "Yes", "No" to 1,0
```

```
geom_line(aes(x_new, p_hat), colour = "blue") +
ylab("Probability of Defaulting")
```



In general, a GLM is three pieces:

1. The random component probability distribution from exponestid family

2. The systemic component A link function relating the parameter of interest (mean!) to
$$\theta$$

$$E[y] = \overline{g}'(\eta)$$

3. A linear predictor

Yin Benoulli (pi)

$$p_{i}^{*} = \frac{\exp(\theta_{i})}{1 + \exp(\theta_{i})} = \bar{q}^{*}(\theta_{i})$$
Note if $Y \sim \text{Ben}(p)$, $E[Y] = p$.

Explanation: describes generating mechanism of observed data.

transforms linear relationship to be on a scale that waters core for parameter of Intrest " links" linear relationship to men.

A linear relationship describing how O is a liver fruitin of predictors.

- ① Standard finalistical denotes link function by \vec{g} : $p = \vec{q}'(\theta) = \frac{\exp(\theta)}{1+\exp(\theta)}$ $\Rightarrow \theta = g(p) = \log(\frac{p}{1-p})$
- (2) Parameter of intest is still the mean, just like linear regression.
- 3) Theoretical reasons for exponential family ... relationship by param of wheest + Variance.

Example (Poisson regression):

- (1) Y: N Poisson (A;)
- (2) $\lambda_i = \exp(\theta_i) = g^{-1}(\theta_i)$ $\theta_i = g(\lambda_i) = \log(\lambda_i)$ | log link"
- 3 θ: = xi β.