

GLM

Questions

- Stormboard: ID: 664949 Key: yellow25

Binomial Regression

Logistic -> Binomial

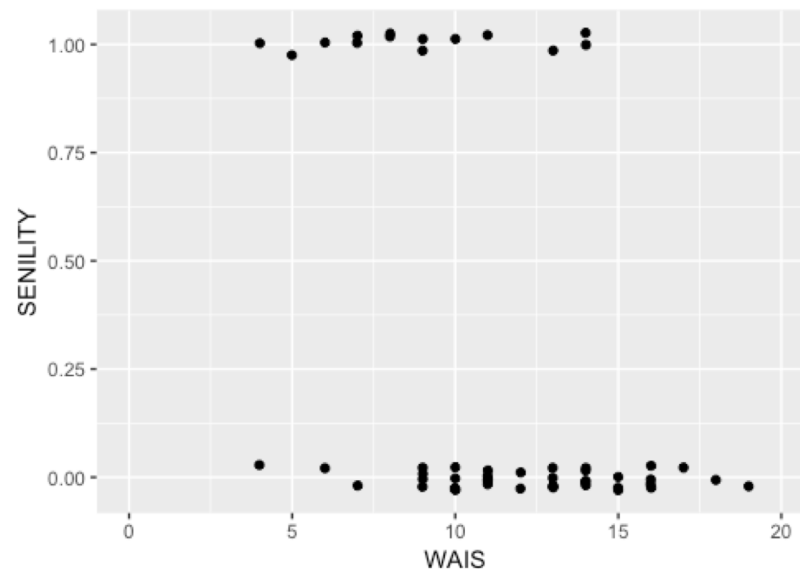
- Bernoulli (probability of one coin flip)
- Binomial (probability of n coin flips)

$$f(y_i, \mu) = \binom{n_i}{y_i} \pi_i^{y_i} (1 - \pi_i)^{n_i - y_i}$$

Senility and Wechsler Adult Intelligence Scale

- 54 elderly people
 - Psychiatric examination to determine signs of senility (0/1).
 - Also measured subset of Wechsler Adult Intelligence Scale (WAIS) (0-20)

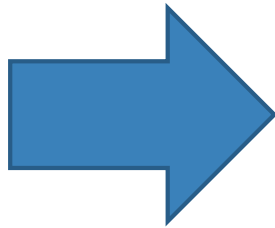
	WAIS	SENILITY
1:	6	0
2:	8	1
3:	13	1
4:	15	0
5:	4	0
6:	12	0
7:	10	0
8:	15	0
9:	7	1
10:	18	0



Senility and Wechsler Adult Intelligence Scale

- 54 elderly people,
 - Psychiatric examination to determine signs of senility.
 - Also measured subset of Wechsler Adult Intelligence Scale (WAIS)

	WAIS	SENILITY
1:	6	0
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4:	15	0
5:	4	0
6:	12	0
7:	10	0
8:	15	0
9:	7	1
10:	18	0
.		



Logistic regression

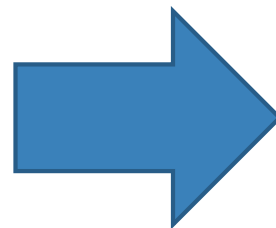
```
glm(formula = SENILITY ~ WAIS, family = binomial,
data = swdt)

            coef.est coef.se
(Intercept)   2.40     1.19
WAIS          -0.32     0.11
---
n = 54, k = 2
residual deviance = 51.0, null deviance = 61.8
(difference = 10.8)
```

Aggregation

- n=54 elderly people can be grouped by G=17 unique WAIS groups

	WAIS	SENILITY
1:	6	0
2:	8	1
3:	13	1
4:	15	0
5:	4	0
6:	12	0
7:	10	0
8:	15	0
9:	7	1
10:	18	0

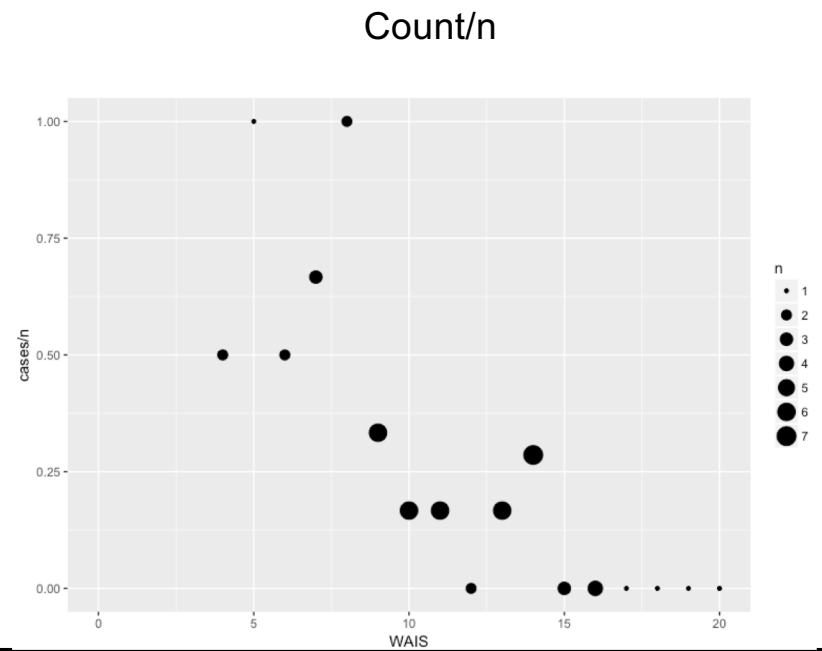
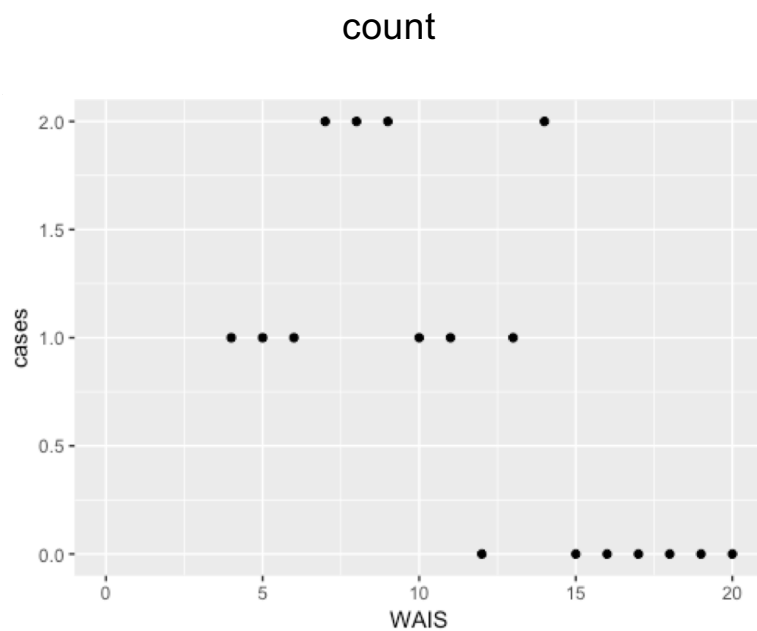


```
swdt%>%group_by(WAIS)%>%  
summarise(cases=sum(SENILITY),n=length(SENILITY))
```

```
# A tibble: 17 x 3  
  WAIS cases    n  
  <int> <int> <int>  
1     4     1     2  
2     5     1     1  
3     6     1     2  
4     7     2     3  
5     8     2     2  
6     9     2     6  
7    10     1     6  
8    11     1     6  
9    12     0     2  
10   13     1     6  
11   14     2     7
```

Count or proportion?

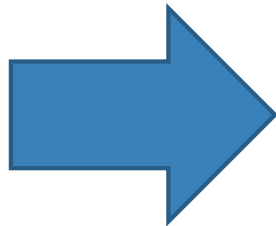
- Key distinction is are these numbers capped?



Aggregation

- n=54 elderly people can be grouped by G=17 unique WAIS groups

```
# A tibble: 17 x 3
  WAIS cases    n
  <int> <int> <int>
1     4     1     2
2     5     1     1
3     6     1     2
4     7     2     3
5     8     2     2
6     9     2     6
7    10     1     6
8    11     1     6
9    12     0     2
10   13     1     6
11   14     2     7
```



Binomial logistic regression

```
glm(formula = cbind(cases, n - cases) ~ WAIS,
     family = binomial,
     data = swdtsum)
               coef.est coef.se
(Intercept)   2.40      1.19
WAIS          -0.32      0.11
---
n = 17, k = 2
residual deviance = 9.4, null deviance = 20.2
(difference = 10.8)
```

Compare the result

- Logistic

```
glm(formula = SENILITY ~ WAIS, family = binomial,
data = swdt)

              coef.est coef.se
(Intercept)   2.40      1.19
WAIS          -0.32      0.11
---
n = 54, k = 2
residual deviance = 51.0, null deviance = 61.8
(difference = 10.8)
```

- Binomial

```
glm(formula = cbind(cases, n - cases) ~ WAIS, family
= binomial, data = swdtsum)

              coef.est coef.se
(Intercept)   2.40      1.19
WAIS          -0.32      0.11
---
n = 17, k = 2
residual deviance = 9.4, null deviance = 20.2
(difference = 10.8)
```

Displaying the fitted curve

```
glm(formula = SENILITY ~ WAIS, family = binomial,  
data = swdt)
```

	coef.est	coef.se
(Intercept)	2.40	1.19
WAIS	-0.32	0.11

n = **54**, k = 2

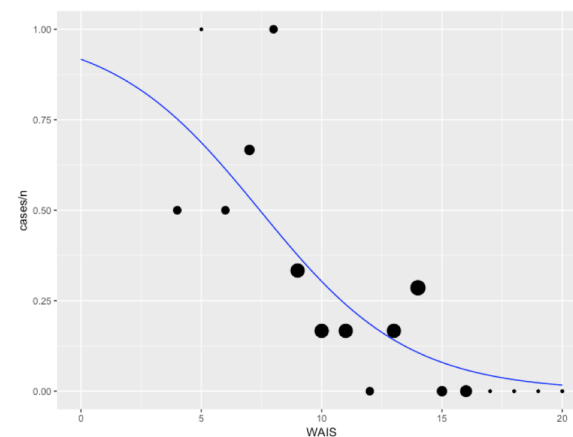
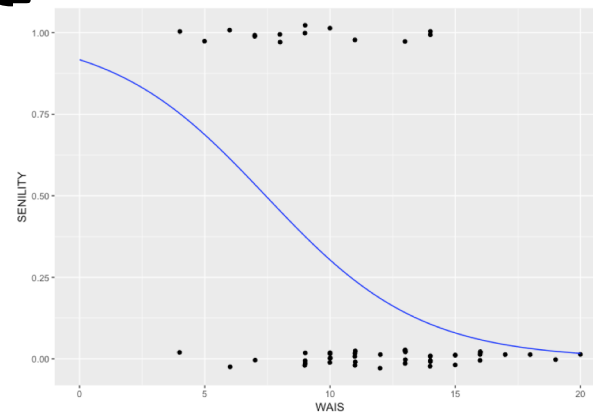
residual deviance = **51.0**, null deviance = **61.8**
(difference = 10.8)

```
glm(formula = cbind(cases, n - cases) ~ WAIS, family  
= binomial, data = swdtsum)
```

	coef.est	coef.se
(Intercept)	2.40	1.19
WAIS	-0.32	0.11

n = **17**, k = 2

residual deviance = **9.4**, null deviance = **20.2**
(difference = 10.8)



Chisquare goodness of fit test

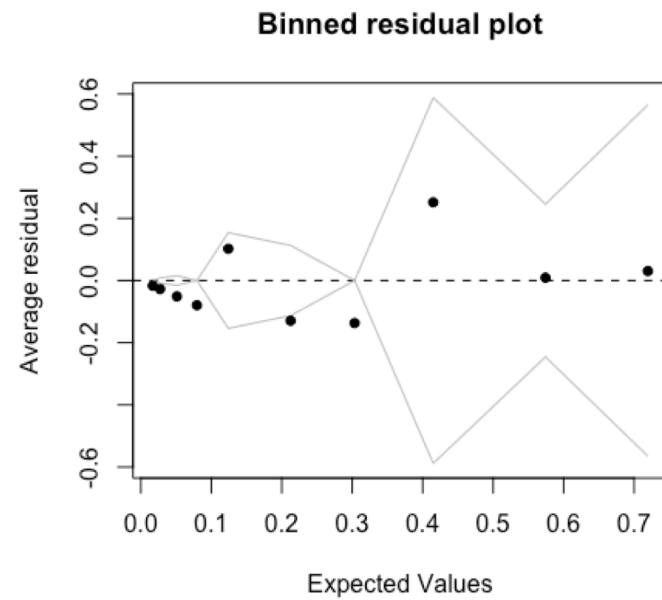
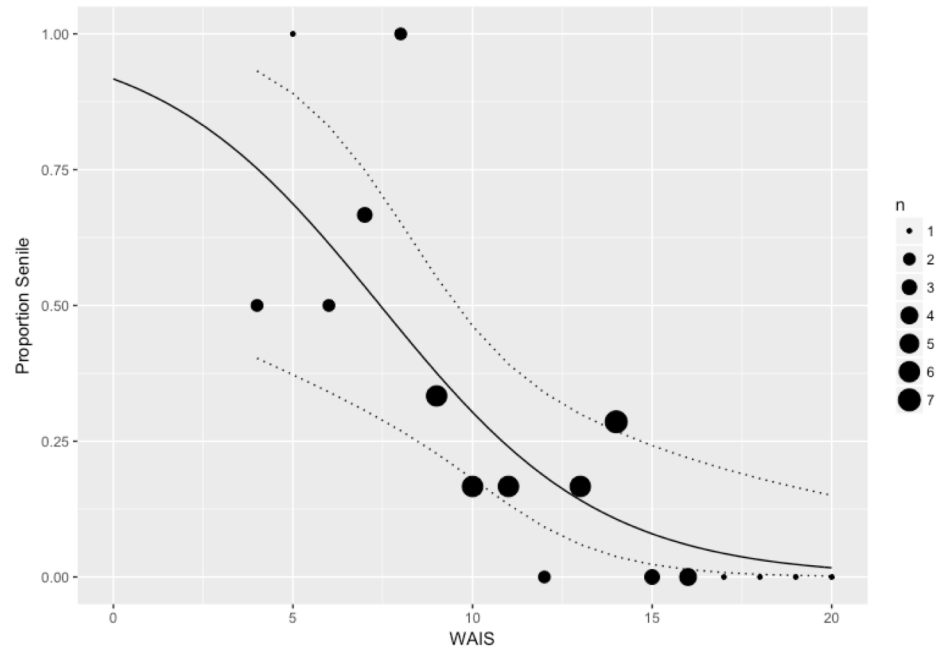
- Should be identical for the two

```
> with(bernlogitfit, pchisq(null.deviance - deviance,  
df.null - df.residual, lower.tail = FALSE))  
[1] 0.001021086  
> with(binlogitfit, pchisq(null.deviance - deviance,  
df.null - df.residual, lower.tail = FALSE))  
[1] 0.001021086
```

AIC

```
> AIC(bernlogitfit)
[1] 55.01738
> AIC(binlogitfit)
[1] 27.79186
```

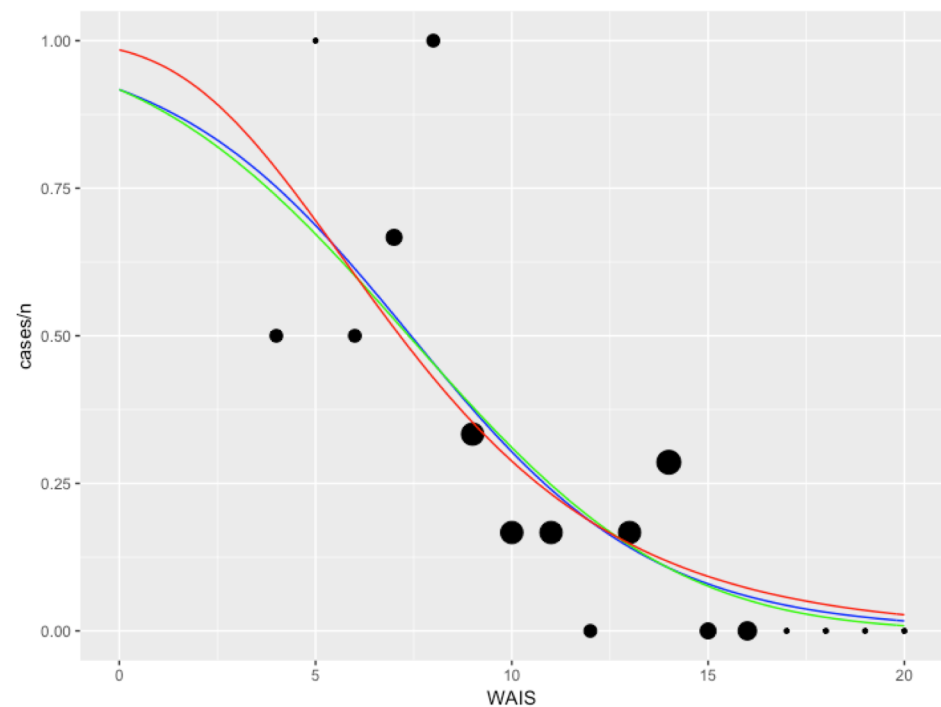
Checking the fit



Different link functions

- Blue logistic
- Green: probit
- Red: c-log-log

```
> AIC(binlogitfit)
[1] 27.79186
> AIC(binprobitfit)
[1] 27.75807
> AIC(bincloglogfit)
[1] 28.08519
```



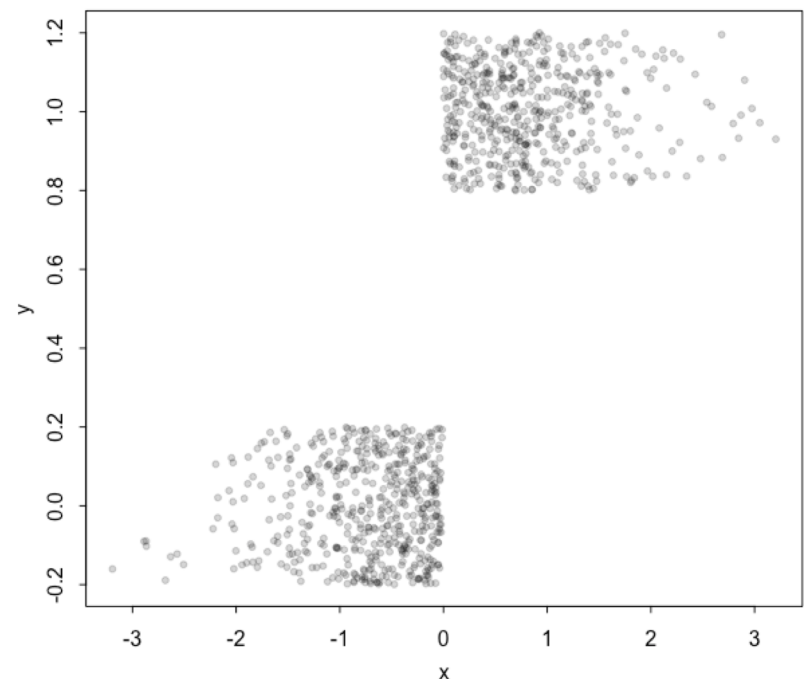
Be careful with the warnings!!

- When you have linear separation

```
> x<-rnorm(1000)
> y<-1*(x>0)
>
> binlogitfit<-glm(y~x,
data=data.frame(x,y),family=binomial)
```

Warning messages:

1: glm.fit: algorithm did not converge
2: glm.fit: fitted probabilities numerically 0 or 1 occurred




```
> summary(binlogitfit)
```

Call:

```
glm(formula = y ~ x, family = binomial, data = data.frame(x, y))
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-0.003301	0.000000	0.000000	0.000000	0.003318

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	10.47	241.68	0.043	0.965
x	3081.20	46607.71	0.066	0.947

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 1.3862e+03 on 999 degrees of freedom
Residual deviance: 2.1910e-05 on 998 degrees of freedom
AIC: 4

Number of Fisher Scoring iterations: 25

| Logistic regression coefficient | > 5 means

- One unit difference can make the probability go from 50:50 to 99%

```
> invlogit(0)
[1] 0.5
> invlogit(0+5)
[1] 0.9933071
> invlogit(0-5)
[1] 0.006692851
```

- If you have such a predictor, why bother with logistic regression?

Logistic or Poisson

- Number of elderly person that was senile out of particular study group.
- Number of senile elderly person that was counted in elderly care facility.
- Number of mouse that survived in toxicity experiment.
- Number of mouse that was counted dead after exterminator sprayed pesticides.
- Number of successful mating after certain number of trials for elephants.
- Number of mating for an elephant.

Poisson Regression

