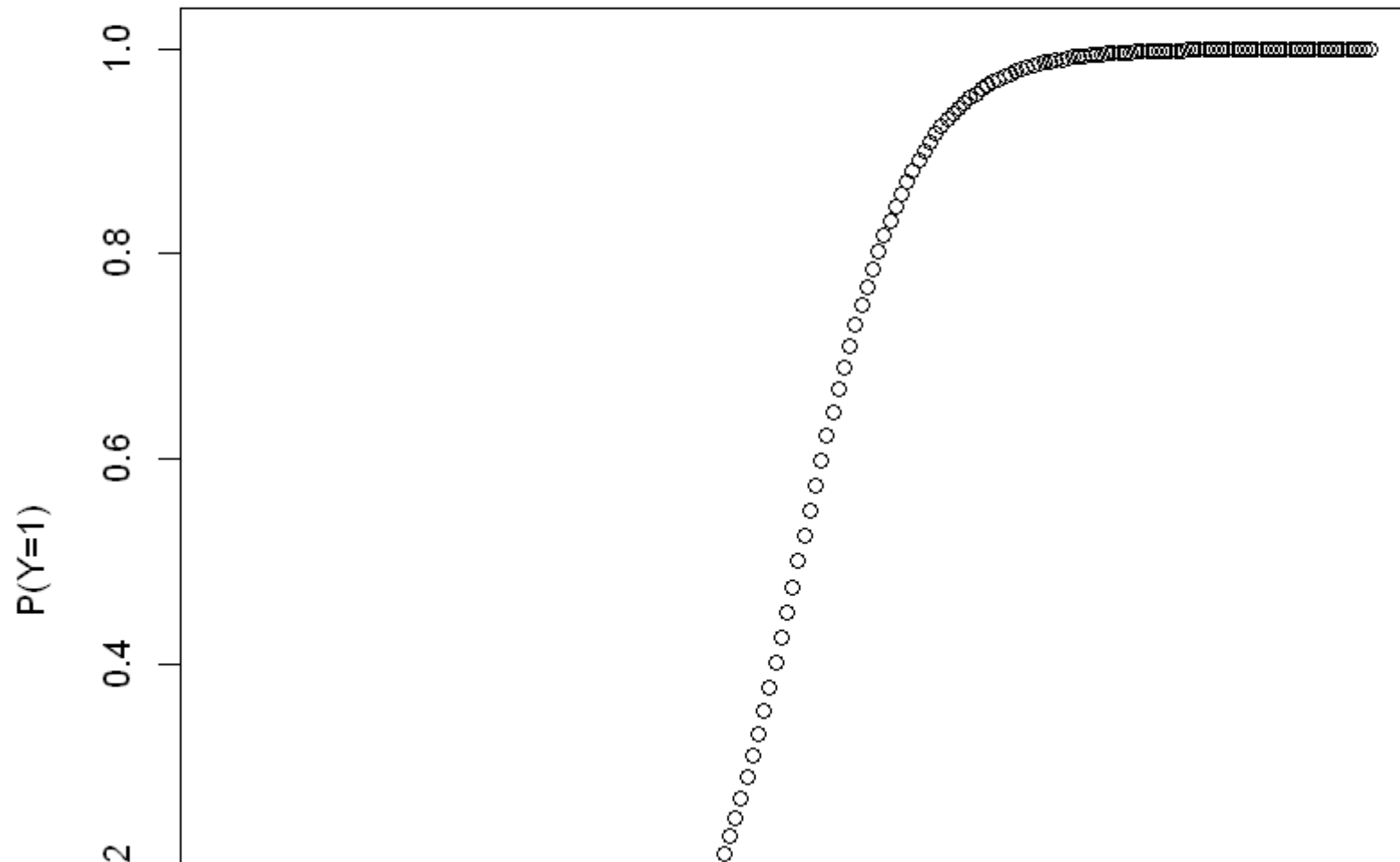
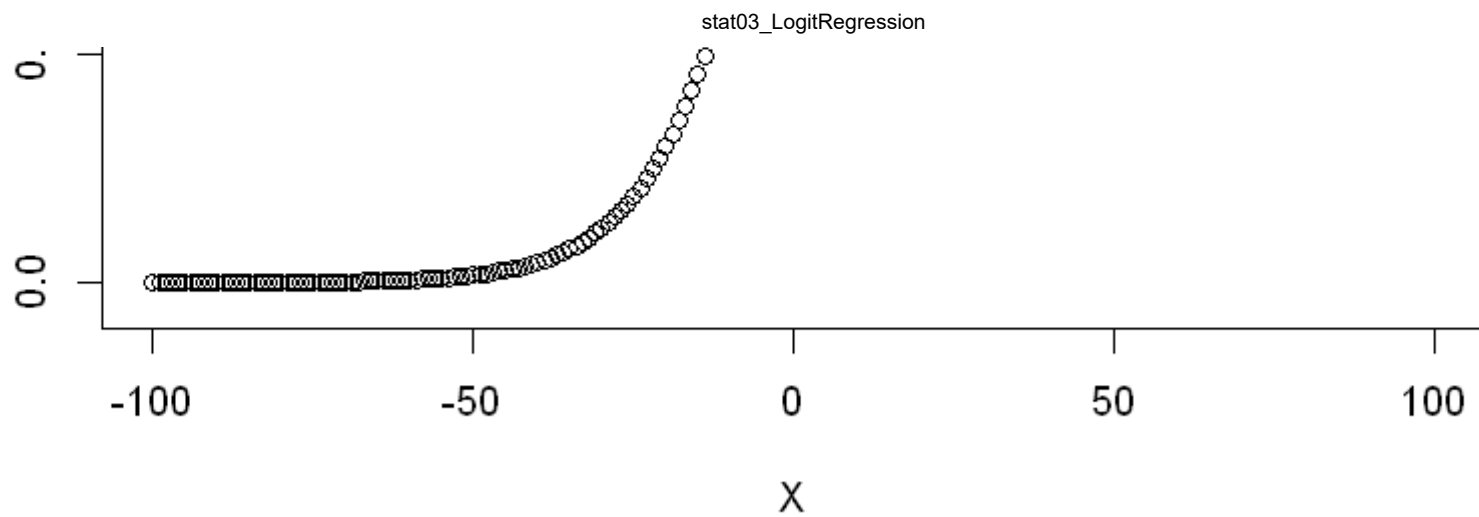


## 로지스틱 회귀 분석 실습

```
In [6]: x <- c(-100:100)
b = 0 # intercept(절편)
m = 0.1 # slope
y = exp((b + m*x)) / (1 + exp((b+m*x)))
plot(x, y, xlab="X", ylab="P(Y=1)")
title(main="Sigmoid (Logistic) Function")
```

## Sigmoid (Logistic) Function





## 우리집 강아지 혈떡거림 - 로지스틱 회귀 모형

여러분의 개가 혈떡거릴때가 있다. 어느 경우에 혈떡일까? 조사해 보니, 온도에 의해 영향을 많이 받는 것 같았다.



<https://www.flickr.com/photos/125021464@N07/18533460483> (<https://www.flickr.com/photos/125021464@N07/18533460483>) (참조)

```
In [7]: temp <- c(10, 13.2, 12.1, 15, 17, 18, 15, 14.9, 16.0, 19,  
                21, 22, 24.5, 28, 27, 20.1, 25.6, 27.2, 29, 28.2)  
dog.panting <- c(0, 0, 0, 0, 0, 1, 0, 1, 0, 0,  
                1, 1, 1, 1, 0, 1, 0, 1, 1, 1)
```

```
In [8]: data <- data.frame(temp, dog.panting)
data
```

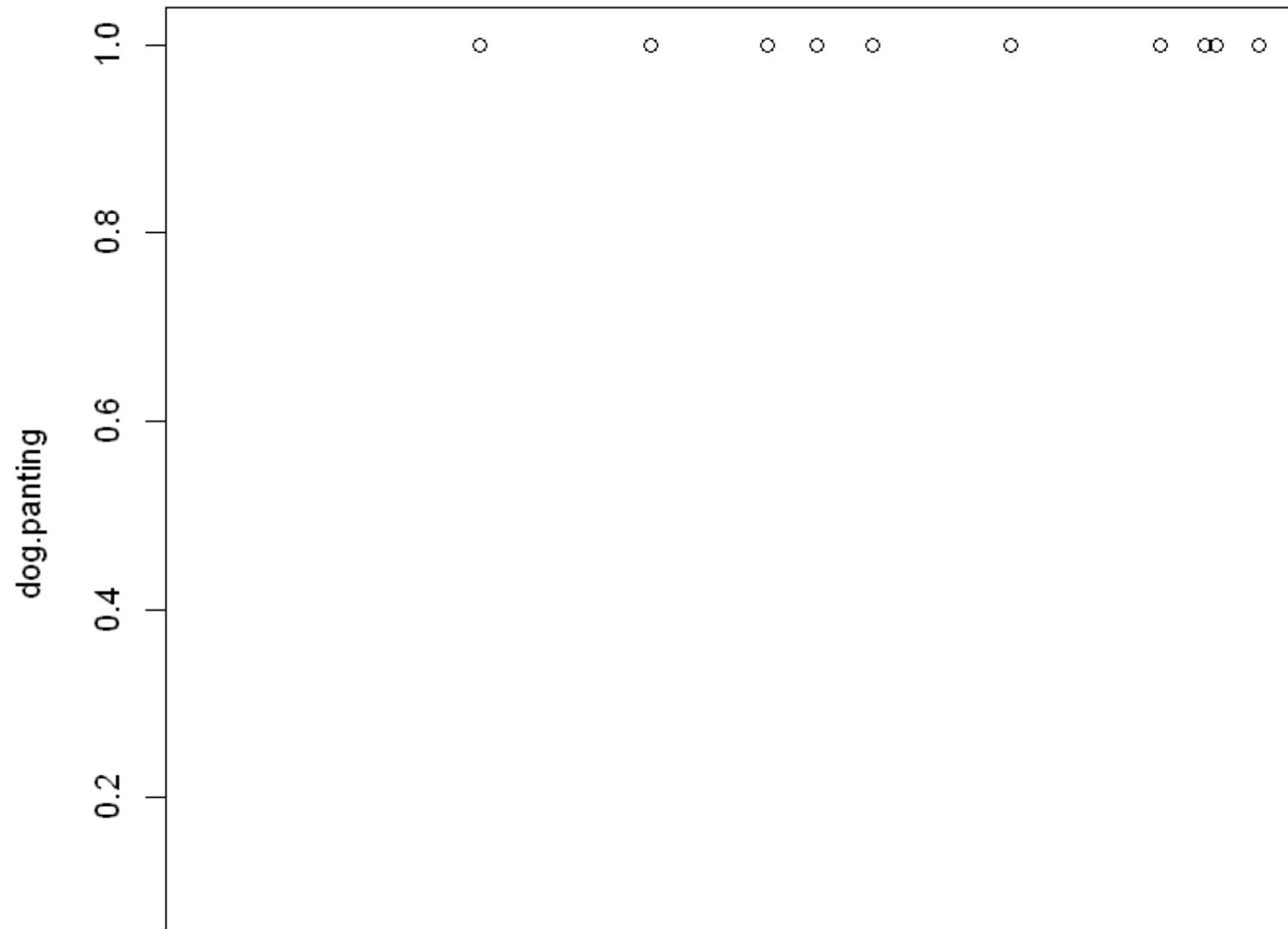
temp	dog.panting
10.0	0
13.2	0
12.1	0
15.0	0
17.0	0
18.0	1
15.0	0
14.9	1
16.0	0
19.0	0
21.0	1
22.0	1
24.5	1
28.0	1
27.0	0
20.1	1
25.6	0
27.2	1
29.0	1
28.2	1

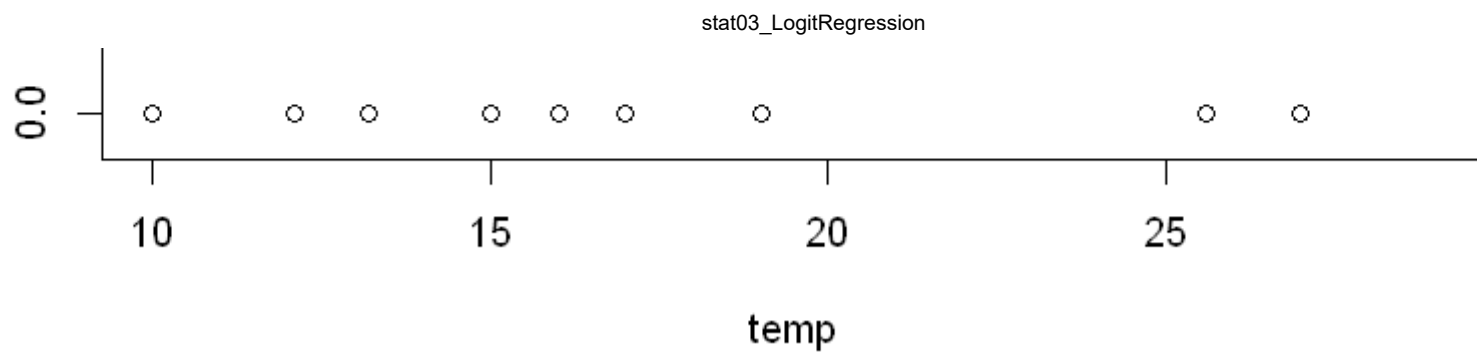
## 데이터 셋 설명

- temp : 온도

- dog.panting (0: 헐떡임, 1: 헐떡이지 않음)

```
In [9]: plot(dog.panting ~ temp)
```





로지스틱 회귀 모델 만들기



```
In [11]: model <- glm(dog.panting~temp, data=data,
                    family = 'binomial')
dim(data)
summary(model)
```

20 2

Call:  
glm(formula = dog.panting ~ temp, family = "binomial", data = data)

Deviance Residuals:

Min	1Q	Median	3Q	Max
-1.87261	-0.76187	0.02904	0.84385	1.69679

Coefficients:

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	-4.5334	2.1439	-2.115	0.0345 *
temp	0.2258	0.1042	2.167	0.0302 *

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 27.726 on 19 degrees of freedom  
Residual deviance: 21.339 on 18 degrees of freedom  
AIC: 25.339

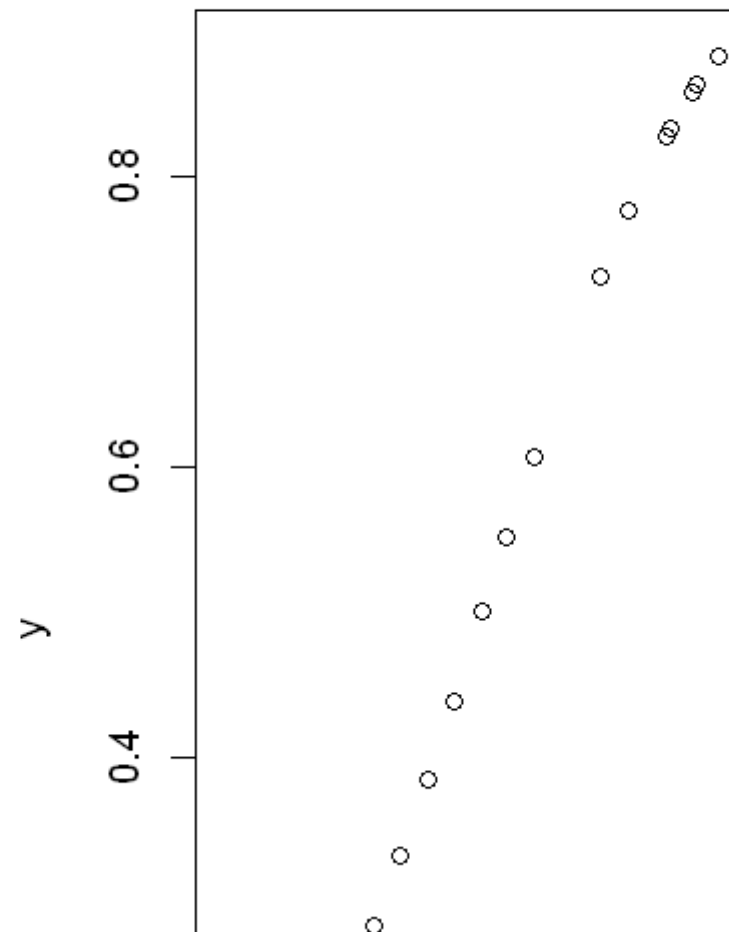
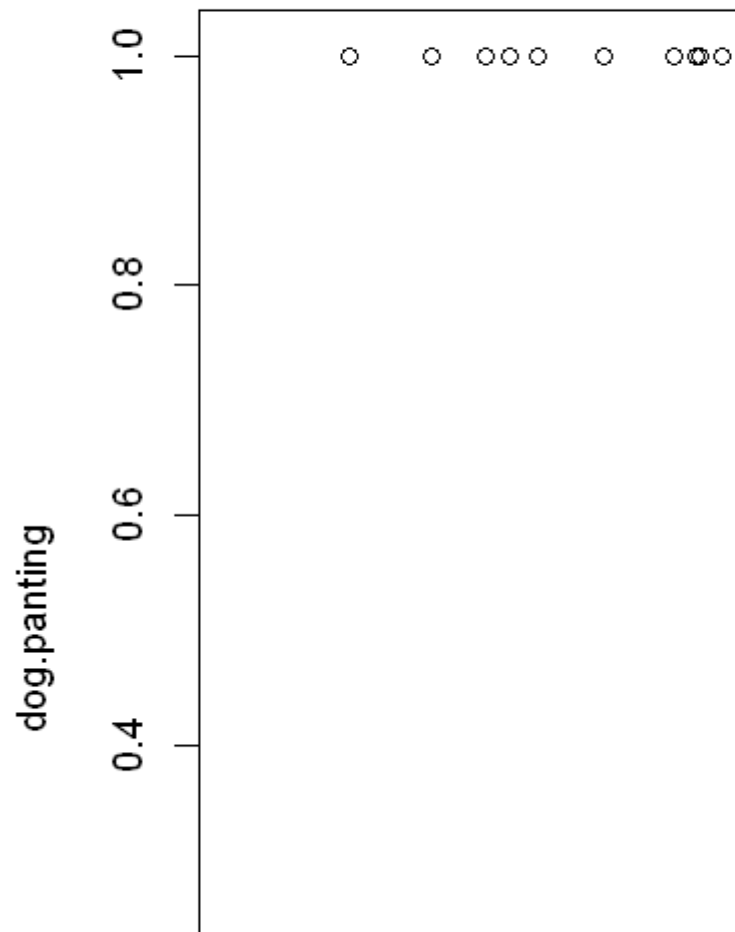
Number of Fisher Scoring iterations: 3

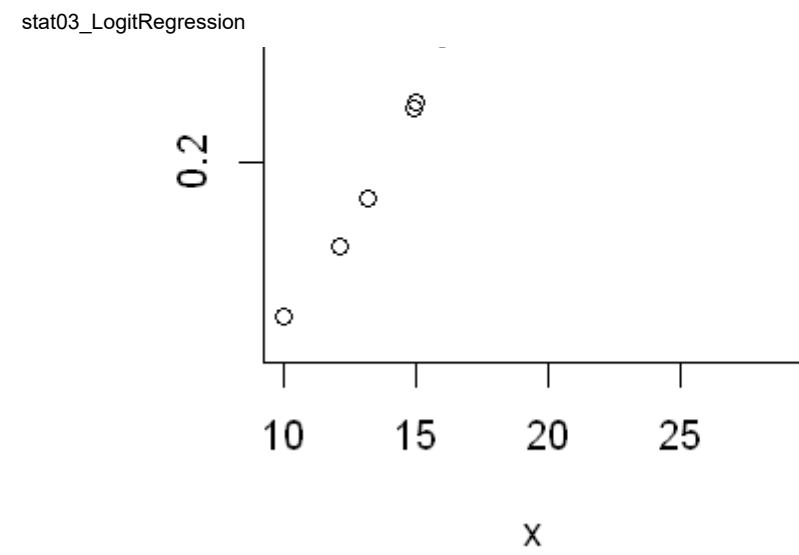
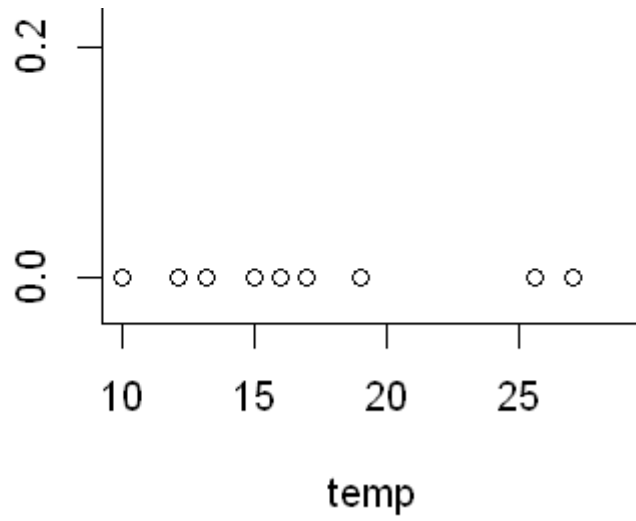
## 예측값과 잔차들 그래프

```
In [12]: model$coefficients
```

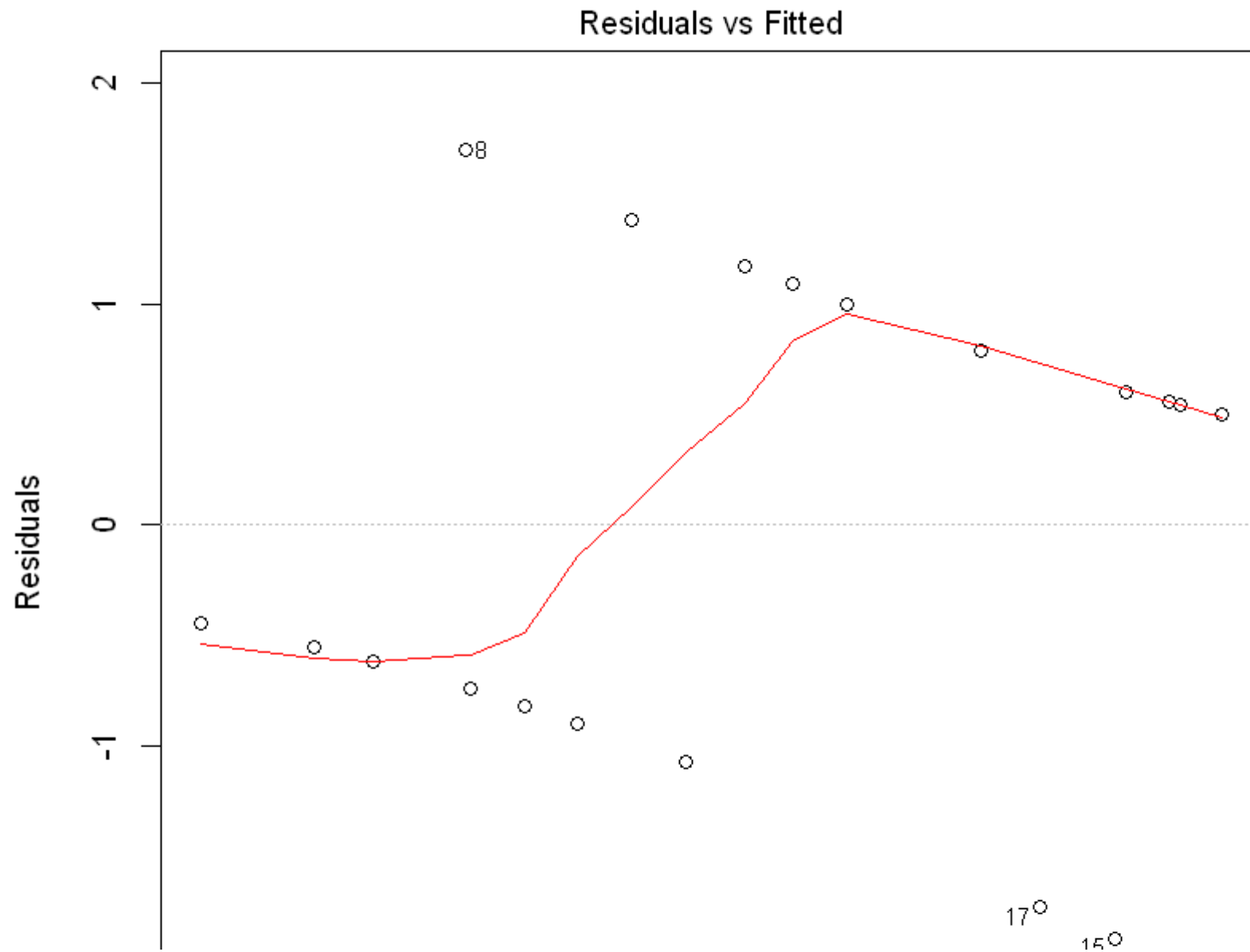
<b>(Intercept)</b>	-4.53338990823194
<b>temp</b>	0.225797708736263

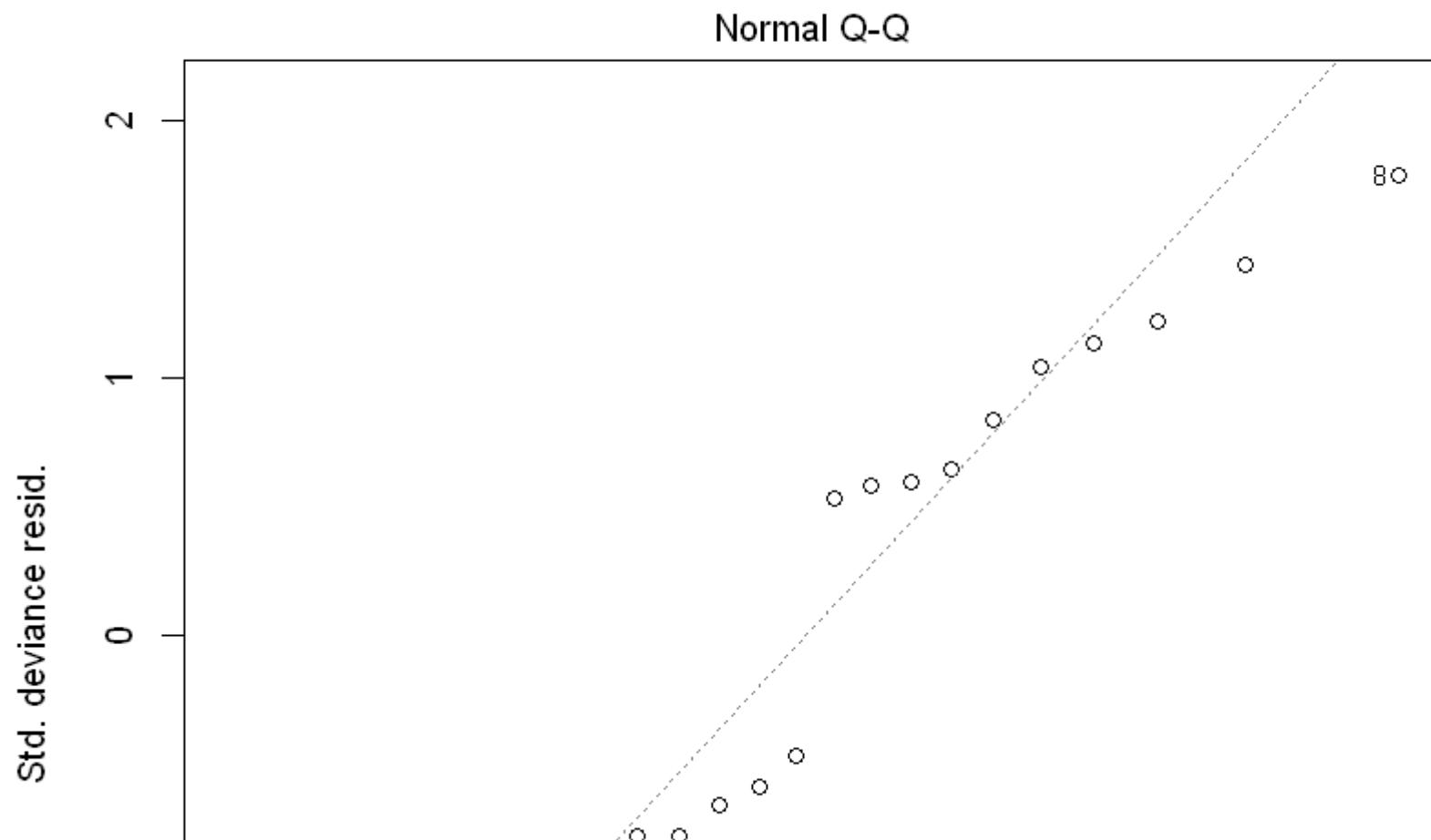
```
In [15]: x <- temp
b <- model$coefficients[1] # intercept
m <- model$coefficients[2] # slope
y = exp((b + m*x)) / (1 + exp((b + m*x)))
par(mfrow=c(1,2))
plot(temp, dog.panting)
plot(x,y)
```

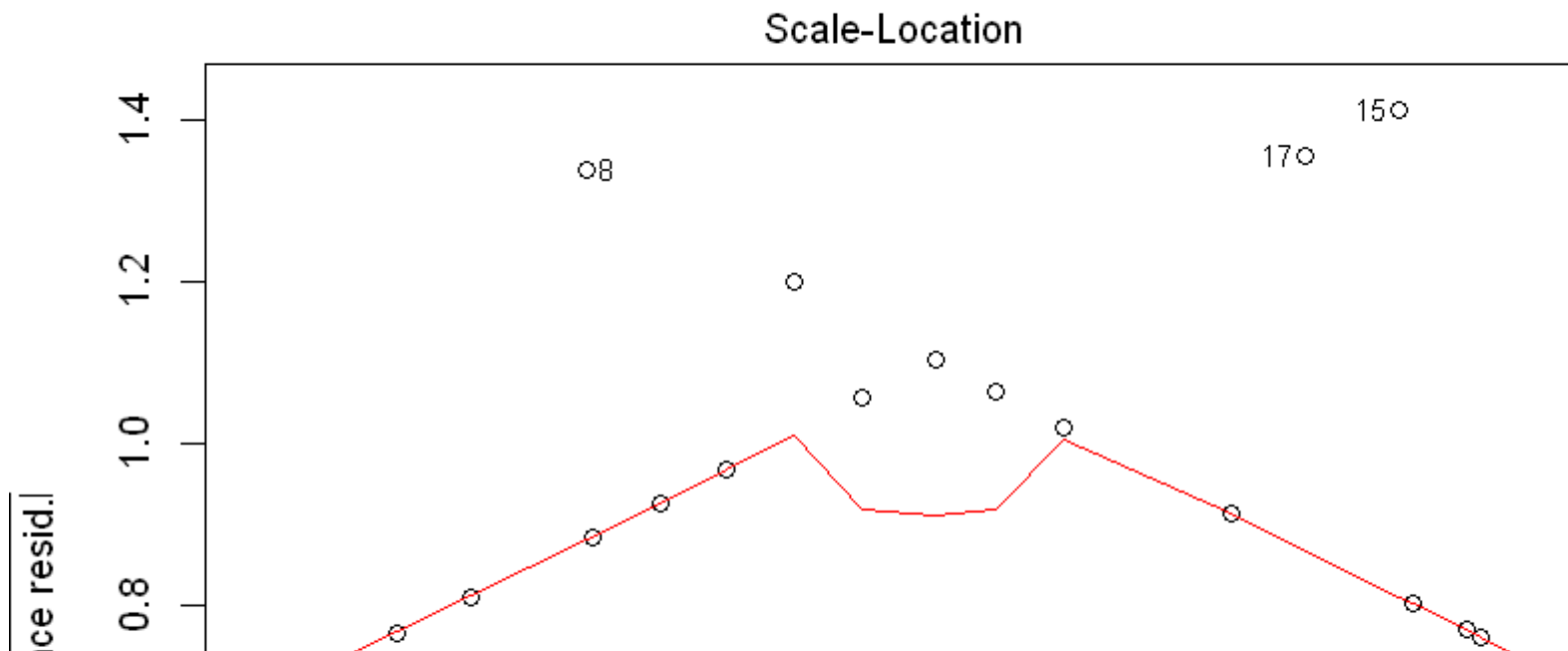
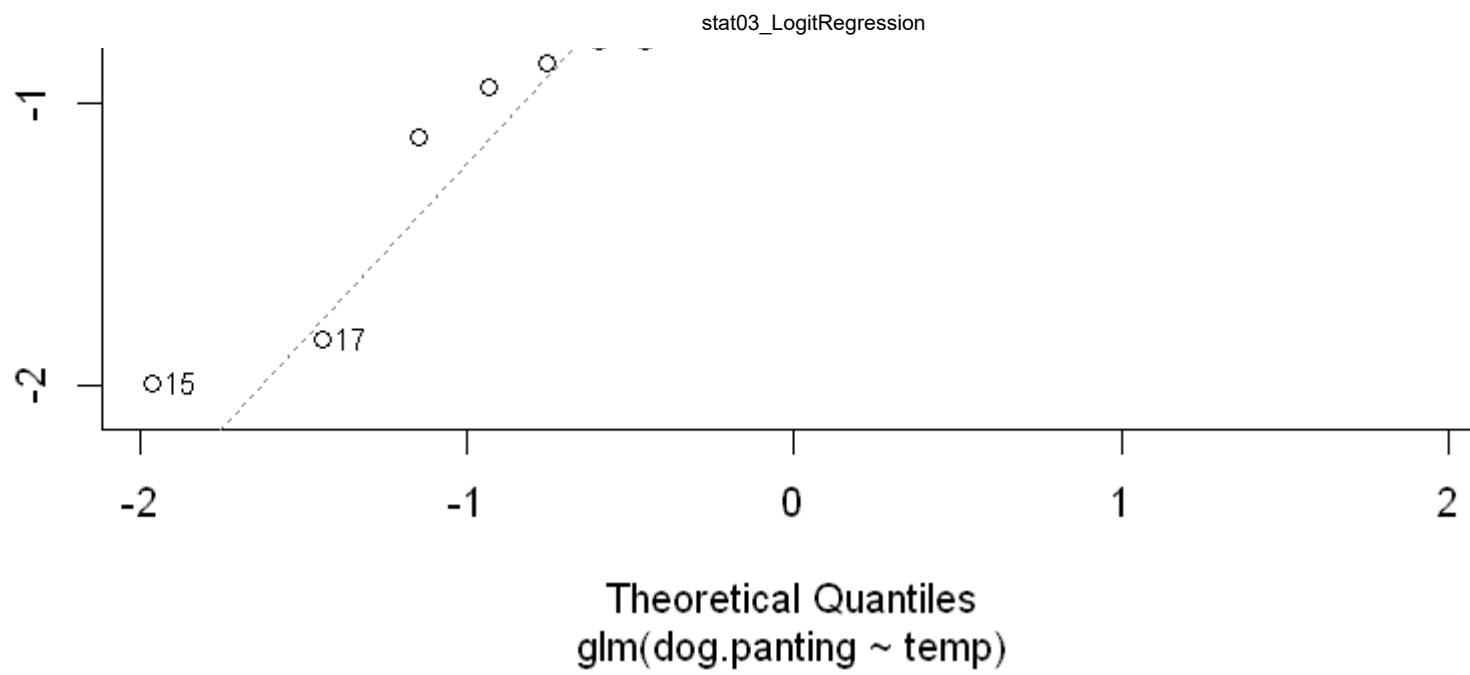


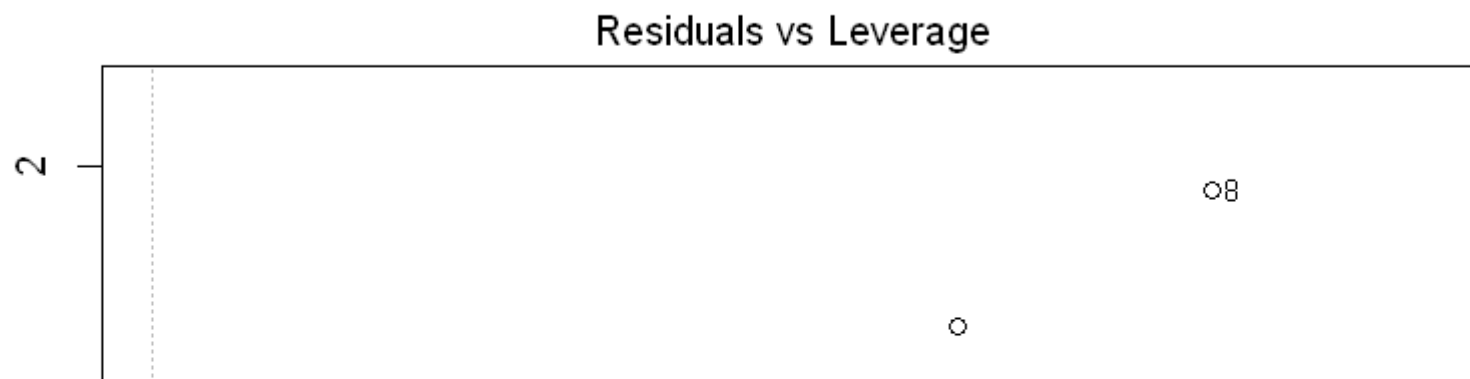
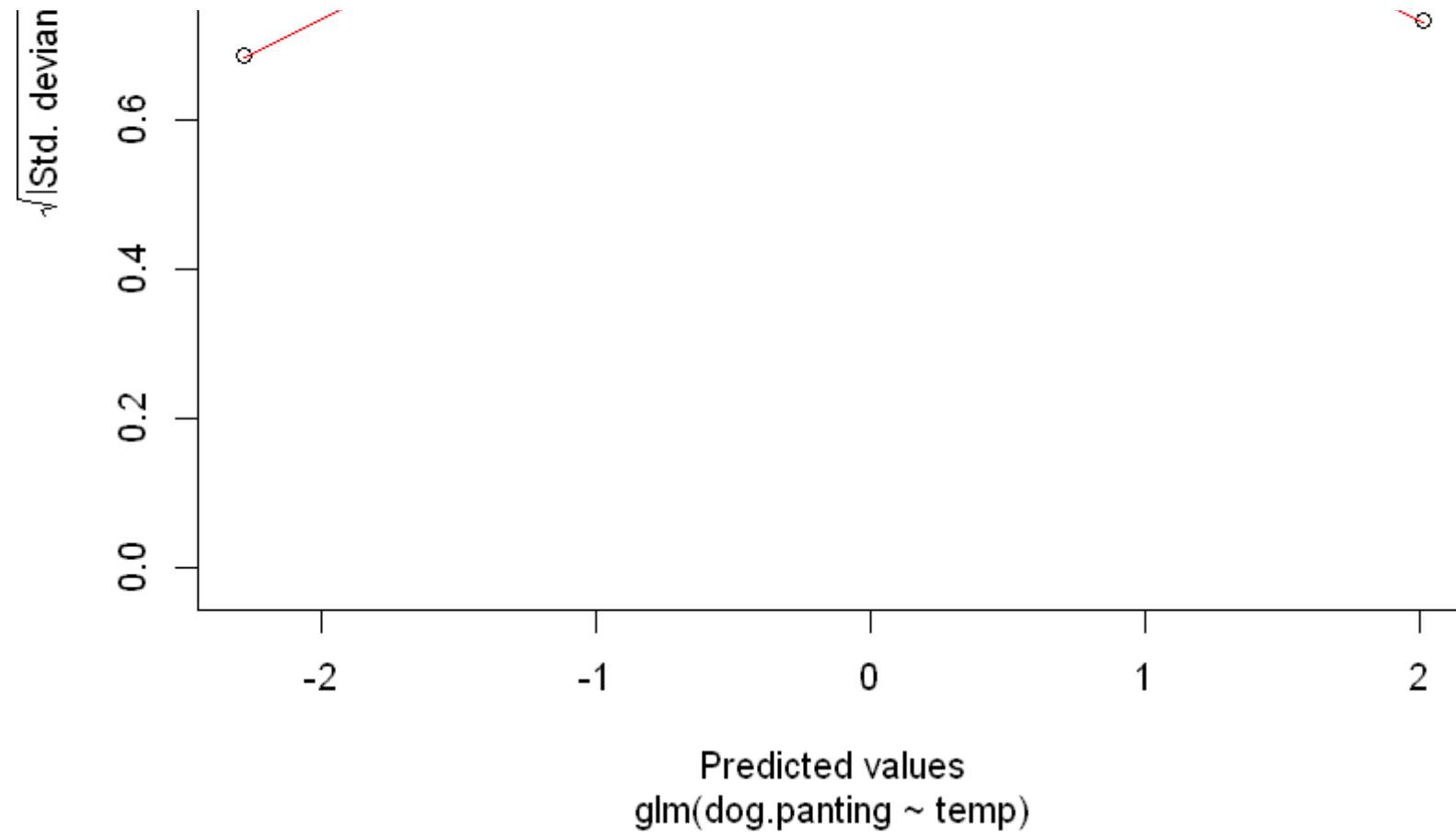


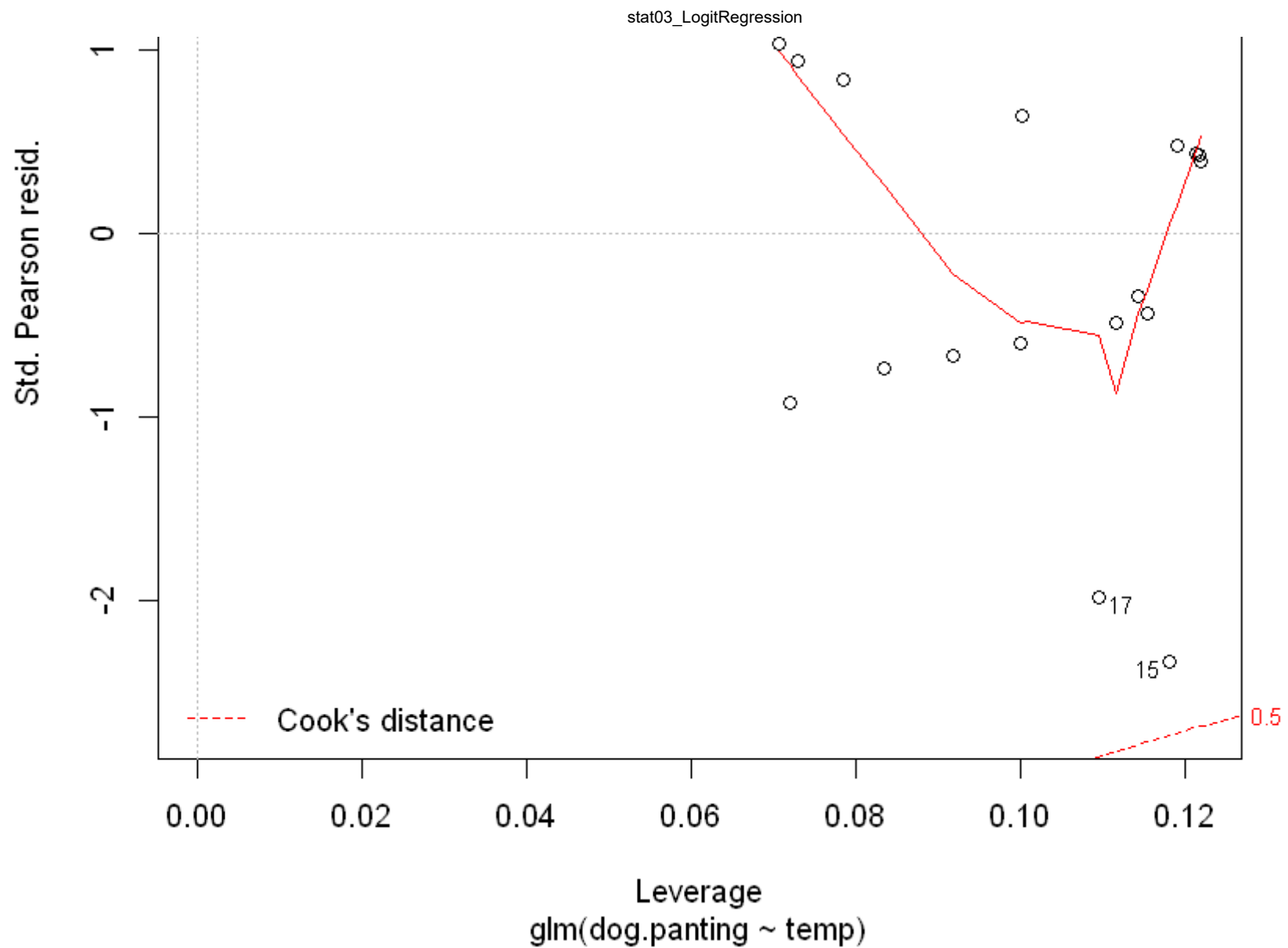
```
In [14]: plot(model)
```











### 로지스틱 회귀 분석 예

- mtcars 데이터 셋은 1974년 Motor Trend US magazine의 데이터 셋이다.
- 32종류의 자동차를 10가지 항목을 조사한 데이터 셋이다.
- 총 10가지 항목은 다음과 같다.



mpg : 연비(Miles/ (US) gallon)  
 cyl : 실린더 개수(Number of cylinders)  
 disp : 배기량 (Displacement (cu.in))  
 hp : 마력(Gross horsepower)  
 drat : 후방차축 비율(Rear axle ratio)  
 wt : 무게(Weight (1,000 lbs))  
 qsec : 1/4 마일에 도달하는데 소요되는 시간(1/4mile time)  
 vs : 엔진(0 = V engine, 1=S engine)  
 am : 변속기(0=자동, 1=수동)  
 gear : 기어 개수(Number of forward gears)  
 carb : 기화기 개수(Number of carburetors)

## 실습 1

- (1) data(mtcars)를 불러온다.
- (2) subset 또는 select, 또는 기본 열 선택을 이용하여 mpg, am, vs 열을 선택한다.
- (3) glm 함수를 이용하여 로지스틱 회귀 분석을 수행해보자. (vs ~ mpg)
- (4) summary ( ) 함수를 이용하여 AIC 통계량을 확인해 보자.

In [17]:

	mpg	am	vs
<b>Mazda RX4</b>	21.0	1	0
<b>Mazda RX4 Wag</b>	21.0	1	0
<b>Datsun 710</b>	22.8	1	1
<b>Hornet 4 Drive</b>	21.4	0	1
<b>Hornet Sportabout</b>	18.7	0	0
<b>Valiant</b>	18.1	0	1

In [19]:

```
Call:
glm(formula = vs ~ mpg, family = "binomial", data = dat)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-2.2127  -0.5121  -0.2276   0.6402   1.6980

Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)  -8.8331     3.1623  -2.793  0.00522 **
mpg           0.4304     0.1584   2.717  0.00659 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

    Null deviance: 43.860  on 31  degrees of freedom
Residual deviance: 25.533  on 30  degrees of freedom
AIC: 29.533

Number of Fisher Scoring iterations: 6
```

In [ ]: