hw3

Giang Vu

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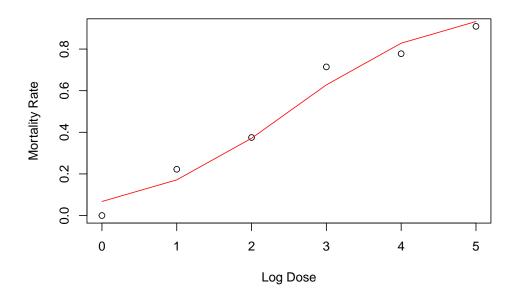
### Homework 3

#### (a)

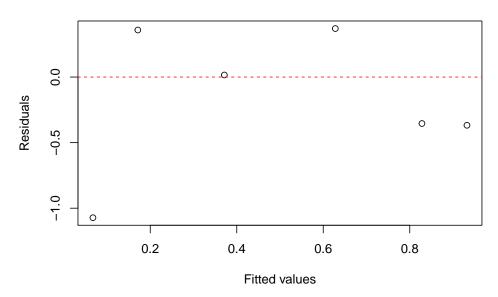
The model I am assuming is  $\begin{aligned} logit(\pi_i) &= \beta_0 + \beta_1 log_2(X_i) \\ \text{where } \pi_i &= Y_i/m \text{ is the mortality rate.} \\ \beta_0 \text{ is log odds of } Y_i &= 1 \text{ when } log_2(X_i) = 0. \\ \beta_1 \text{ is the log odds ratio between log dose } X_{i+1} \text{ and log dose } X_i. \\ Y_i | X_i \text{ are assumed to be independent.} \end{aligned}$ 

### (b)

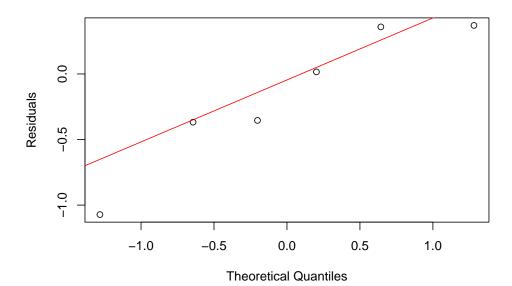
After fitting the model using quasibinomial family with logic link, the plot of the fitted mortality rate and raw mortality fractions y/m against log dose is obtained below. Looking at the fit of the model, we could see that it fits quite well. I checked residual plot and qq-plot and the constant variances and normality assumptions are both satisfied as well.



## Residual plot



### Normal Q-Q Plot



(i) 
$$p = LD_{50} \Rightarrow logit(\pi) = \hat{\beta}_0 + \hat{\beta}_1 log_2(\pi) = 0$$

$$\hat{\gamma} = 2 \left( \frac{\hat{\beta}_0}{\hat{\beta}_1} \right) = 2 \left( \frac{2.626}{1.050} \right) \approx 5.66$$

An Estimate for  $\theta'$  is 5.66

(ii) We want to test null hypothesis

Ho:  $log_2(T) = \frac{\beta_0}{\beta_1} = 4$ 

$$\Rightarrow \beta_0 = 4\beta_1$$
Our restricted model is  $logit(\pi_i) = -4\beta_1 + \beta_1 log_2(\chi_i)$ 

$$= ) logit(\pi_i) = \beta_1 (log_2(\chi_i) - 4)$$

After filling, this model, we obtain coefficient estimate
$$\hat{\beta}_1 = 0.5528$$
After doing the lihood vatrotest, the likelihood vatrotest  $\theta$  with  $\theta$ -value of  $\theta$ . If Ho is true,  $\theta$ . In (4)  $\theta$   $\theta$   $\theta$ .

We fail to reject to  $\theta$ . If Ho is true,  $\theta$ .

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(c)

(iii) I didn't understand what "plot the restricted log likelihood against the hypothesized value of log2(gamma)" is, because restricted log likelihood is one number, and the hypothesized value of log2(gamma) is 4 (null hypothesis in the previous part), so if I were to plot it it would be just a single point in the plane, so I did not understand how that would help with constructing a confidence interval. I used some built in function to generate the 95% confidence interval for the parameter of the restricted model to be (0.083, 1.14), so the 95% confidence interval for gamma would be (1.06, 2.20). This also doesn't really make sense because we fail to reject that gamma = 4 earlier, but 4 is outside of this confidence interval.

cl) (i) 
$$\hat{B}^{2} = \frac{1}{|Y|-2} \sum_{x \in Y} \frac{\{y(2) - m(x) \hat{\Pi}(x)\}^{2}}{m(x) \hat{\Pi}(x) \{1 - \hat{\Pi}(x)\}^{2}}$$

=)  $\hat{B}^{2} (1TI-2) = \sum_{x \in Y} \frac{\{y(x) - m(x) \hat{\Pi}(x)\}^{2}}{m(x) \hat{\Pi}(x) \{1 - \hat{\Pi}(x)\}^{2}}$ 

=  $\sum_{x \in Y} \frac{\{y(x) - m(x) \hat{\Pi}(x)\}^{2} - m(x)}{m(x) \hat{\Pi}(x) \{1 - \hat{\Pi}(x)\}^{2}} = \sum_{x \in Y} \frac{\{y(x) - m(x) \hat{\Pi}(x)\}^{2} - m(x)}{m(x) \hat{\Pi}(x) \{1 - \hat{\Pi}(x)\}^{2}} = X^{2}$ 

(ii) Asymptotically,  $X^{2} \sim \chi_{C-P}^{2}$ 

=)  $\chi^{2} \sim Gamma \left(\frac{c-P}{2}, \frac{1}{2}\right)$ 

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