Homework 3: Categorical Data

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Your assignment should be submitted in two separate files by 8am on Thursday October 19th. The first, should be an RMarkdown (.Rmd) or another format that dynamically compiles your write up and runs the code inside it. The second should be the PDF file that was reproducibly compiled using the first file. All figures should be generated by the code, none should be loaded directly. The homework files should be submitted using your shared Google Drive folder with the instructor.

GLMs

Question 1

Write your own R code for obtaining the MLEs and SEs for a Poisson GLM using two covariates. Conduct a small simulation study designed to stress-test your code and show that it can work in a variety of situations. Compare the results to the glm() function in R. Summarize the results and your experience in 1-2 paragraphs. [Acknowledgments to Ciprian Crainiceanu for this question.]

Question 2

CDA exercise 4.11

Table 4 is based on a study with Brithish doctors. (a) For each age, find the sample coronary death rates per 1000 person-years for nonsmokers and smokers. To compare them, take their ratio and describe its dependence on age.

- (b) Fit a main-effects model for the log rates using age and smoking as factors. In discussing lack of fit, show that this model assumes a constant ratio of nonsmokers' to smokers' coronary death rates over age.
- (c) From part (a), explain why it is sensible to add a quantitative interaction of age and smoking. For this model, show that the log ratio of coronary death rates changes linearly with age. Assign scores to age, fit the model, and interpret.

Question 3

CDA exercise 4.12

Table 1: 4.9 Data for Question 6 on Coronary Death Rates

| | Person-y | years | Coronary Deaths | | | | |
|-------|------------|---------|-----------------|---------|--|--|--|
| Age | Nonsmokers | Smokers | Nonsmokers | Smokers | | | |
| 35-44 | 18,793 | 52,407 | 2 | 32 | | | |
| 45-54 | 10,673 | 43,248 | 12 | 104 | | | |
| 55-64 | 5710 | 28,612 | 28 | 206 | | | |
| 65-74 | 2585 | 12,663 | 28 | 186 | | | |
| 75-84 | 1462 | 5317 | 31 | 102 | | | |

Table 2: Q7

| | 100 | 10 2. (| ٠. | | | | | | |
|------------------|-----------|---|--|--|--|--|--|--|--|
| | Histology | | | | | | | | |
| Disease Stage | Ι | | II | | | III | | | |
| | 1 | 2 | 3 | 1 | 2 | 3 | 1 | 2 | 3 |
| | 9 | 12 | 42 | 5 | 4 | 28 | 1 | 1 | 19 |
| | (157 | 134 | 212 | 77 | 71 | 130 | 21 | 22 | 101) |
| | 2 | 7 | 26 | 2 | 3 | 19 | 1 | 1 | 11 |
| | (139 | 110 | 136 | 68 | 63 | 72 | 17 | 18 | 63) |
| | 9 | 5 | 12 | 3 | 5 | 10 | 1 | 3 | 7 |
| | (126 | 96 | 90 | 63 | 58 | 42 | 14 | 14 | 43) |
| | 10 | 10 | 10 | 2 | 4 | 5 | 1 | 1 | 6 |
| | (102 | 86 | 64 | 55 | 42 | 21 | 12 | 10 | 32) |
| | 1 | 4 | 5 | 2 | 2 | 0 | 0 | 0 | 3 |
| | (88 | 66 | 47 | 50 | 35 | 14 | 10 | 8 | 21) |
| | 3 | 3 | 4 | 2 | 1 | 3 | 1 | 0 | 3 |
| | (82 | 59 | 39 | 45 | 32 | 13 | 8 | 8 | 14) |
| | 1 | 4 | 1 | 2 | 4 | 2 | 0 | 2 | 3 |
| | (76 | 51 | 29 | 42 | 28 | 7 | 6 | 6 | 10) |
| | | Disease Stage 1 9 (157 2 (139 9 (126 10 (102 1 (88 3 (82 1 | Disease Stage I 1 2 9 12 (157 134 2 7 (139 110 9 5 (126 96 10 10 (102 86 1 4 (88 66 3 3 (82 59 1 4 | Stage 1 2 3 9 12 42 (157 134 212 2 7 26 (139 110 136 9 5 12 (126 96 90 10 10 10 (102 86 64 1 4 5 (88 66 47 3 3 4 (82 59 39 1 4 1 | Disease Stage I Historian 1 2 3 1 9 12 42 5 (157 134 212 77 2 7 26 2 (139 110 136 68 9 5 12 3 (126 96 90 63 10 10 10 2 (102 86 64 55 1 4 5 2 (88 66 47 50 3 3 4 2 (82 59 39 45 1 4 1 2 | Disease Stage I Histology 1 2 3 1 2 9 12 42 5 4 (157 134 212 77 71 2 7 26 2 3 (139 110 136 68 63 9 5 12 3 5 (126 96 90 63 58 10 10 10 2 4 (102 86 64 55 42 1 4 5 2 2 (88 66 47 50 35 3 3 4 2 1 (82 59 39 45 32 1 4 1 2 4 | Disease Stage I II II 1 2 3 1 2 3 9 12 42 5 4 28 (157 134 212 77 71 130 2 7 26 2 3 19 (139 110 136 68 63 72 9 5 12 3 5 10 (126 96 90 63 58 42 10 10 10 2 4 5 (102 86 64 55 42 21 1 4 5 2 2 0 (88 66 47 50 35 14 3 3 4 2 1 3 (82 59 39 45 32 13 1 4 1 2 4 2 | Disease Stage I Histology 1 II 9 12 42 5 4 28 1 (157 134 212 77 71 130 21 (139 110 136 68 63 72 17 9 5 12 3 5 10 1 9 5 12 3 5 10 1 10 10 10 2 4 5 1 10 10 10 2 4 5 1 10 4 5 2 2 0 0 10 10 2 4 5 1 1 4 5 2 2 0 0 (88 66 47 50 35 14 10 (82 59 39 45 32 13 8 1 | Disease Stage I II III IIII III III IIII III III IIII IIIIII IIII IIII IIII IIII IIII IIII IIII IIII IIII IIIII IIIII IIIII IIIII IIIII IIII IIIII IIIIII< |

Table 4.10 describes survival for 539 males diagnosed with lung cancer. The prognostic factors are histology (H) and state (S) of disease. The assumption of a constant rate over time is often not sensible, and this study divided the time scale (T) into two-month intervals and let the rate vary by the time interval. Let μ_{ijk} denote the expected number of deaths and t_{ijk} the total time at risk for histology i and state of disease j, in follow-up time interval k. Analyses suggested a lack of interaction between T and either prognostic factor (i.e., such proportional hazards models have the same effects of H and S for each time interval).

(a) The main effects model

$$log(\mu_{ijk}/t_{ijk}) = \alpha + \beta_i^H + \beta_j^S + \beta_k^T$$

has deviance 43.9. Explain why df = 52. Does the model seems to fit adequately?

(b) For this model, interpret the estimated effects of S,

$$\hat{\beta}_2^S - \hat{\beta}_1^S = 0.470(SE = 0.174)$$

$$\hat{\beta_3^S} - \hat{\beta_1^S} = 1.324(SE = 0.152)$$

(C) The model that adds an $S \times H$ interaction term has deviance 41.5 with df = 48. Test whether a significantly improved fit results by allowing this interaction.

Question 4

CDA exercise 4.16

For binary data, define a GLM using the log link. Show that effects refer to the relative risk. Why do you think this link is not often used? Hint: What happens if the linear predictor takes a positive value?

Question 5

CDA exercise 4.21

A binomial GLM $\pi_i = \phi(\sum_j \beta_j x_{ij})$ with arbitrary inverse link function ϕ assumes that $n_i Y_i$ has a $bin(n_i, \pi_i)$ distribution. Find w_i , in (4.29) and hence $cov(\hat{\beta})$. For logistic regression, show that $w_i = n_i \pi_i (1 - \pi_i)$.

Question 6

CDA exercise 4.27

For known k, show that the negative binomial distribution (4.13) has exponential family form (4.1) with natural parameter $log[\mu/(\mu+k)]$.