- 1. (a) Since $G^2=46.13>9.49=\chi^2_{4,0.05}$ and $G^2=14.71>7.81=\chi^2_{3,0.05}$, the independence and symmetry models have a definite lack of fit. Since $G^2=3.20<5.99=\chi^2_{2,0.05}$ and $G^2=0.24<3.84=\chi^2_{1,0.05}$, both the ordinal quasi-symmetry and quasi-symmetry models are fit well by the data. Since $G^2=3.20-0.24=2.96<3.84=\chi^2_{1,0.05}$, the QS model does not improve significantly on the OQS model. Thus, the OQS model is the most parsimonious model that adequately fits the data.
 - (b) Since $G^2 = 14.71 0.24 = 14.47 > 5.99 = \chi^2_{2,0.05}$, the QS model significantly improves upon the symmetry model. This implies that marginal homogeneity does not hold for this table.
- 2. (a) \bullet (YU, YN, YR, YA, UNRA)
 - (YUA, YN, YR, UNRA)

Y is conditionally independent of $\{R, N\}$ given $\{U, A\}$. Thus, Y

(b) is also conditionally independent of $\{R\}$ given $\{U, A\}$ and of $\{N\}$ given $\{U, A\}$.



- - (b) We test $H_0: \theta = 1$ versus $H_a: \theta > 1$ using Fisher's exact test. The P-value= 0.0083 and the mid-P-value= 0.0083 0.0074/2 = 0.0046. Both the P-value and the mid-P-value provide strong evidence that a greater proportion of the mice exposed to smoke had tumors than did the control mice.
- 4. (a) Do not reject $H_0: \lambda^{DGE} = 0$ in the saturated model since $G^2 = 1.48 0 = 1.48 < 3.84 = \chi^2_{1,0.05}$. There is insufficient evidence to indicate that the odds ratio between gender and depress differ for the two levels of education.
 - (b) Using the homogeneous association model, reject $H_0: \lambda^{DG}=0$ since $G^2=22.82-1.48=21.34>3.84=\chi^2_{1,0.05}$. There is strong evidence of association between gender and depress, controlling for the two levels of education.
 - (c) All the models except (DG, GE), (DG, DE, GE), and (DGE) have strong evidence of a lack of fit. In part (a), we determined that the three-way interaction was not significant, so we would prefer (DG, DE, GE) to the saturated model. Since $G^2 = 5.39 1.48 = 3.91 > 3.84 = \chi^2_{1,0.05}$, we find that the DE interaction is needed. Thus, we recommend the homogeneous association model, (DG, DE, GE).
- 5. (a) Table I is more appropriate since the data are paired with each individual receiving both allergy tests. To test $H_0: \pi_{1+} = \pi_{+i}$, we use McNemar's test. Since $X^2 = (12-28)^2/(12+28) => 3.84 = \chi^2_{1,0.05}$, we reject H_0 and conclude that the proportion of individuals suffering allergic reactions differs for the two types of penicillin.
 - (b) First, $\widehat{OR} = 28/12 = 2.333$. Then a 95% confidence interval for $\log(OR)$ is $\log(28/12) \pm 1.96\sqrt{1/28+1/12} = 0.847 \pm 0.676$, or (0.171, 1.523). Exponentiate the endpoints to obtain the 95% confidence interval for OR: (1.186, 4.589). The odds of an individual being allergic to type BT penicillin are from 1.2 to 4.6 times the odds of an individual being allergic to Type G penicillin.

1

6. (a) The 95% confidence interval for β_{Tr} is given by $0.9379 \pm (1.96)(0.3310)$ or (0.289, 1.587). We exponentiate the end points to obtain a 95% confidence interval for the OR for the cumulative logit of the treatment group relative to the control group: (1.335, 4.887). Thus, treatment group has a larger cumulative probability $P(y \le j)$ for a fixed age. This implies that the treatment group has a higher probability for low values of y which implies less pain for individuals in this group. Thus, the treatment appears effective.

(b)

$$P[\mathtt{y} \leq 4] = \frac{e^{-3.2271 + 0.9379}}{1 + e^{-3.2271 + 0.9379}} = 0.0902 \Longrightarrow P[\mathtt{y} = 5] = 1 - P[\mathtt{y} \leq 4] = 1 - 0.0902 = 0.9098$$

Note: Due to a data entry error, the intercepts and the coefficient for Age1 were not correct on the output. Intercept 4 should be 2.3825 resulting in $P[y \le 4] = 0.965$ and P[y = 5] = 0.035. This matches up well with the observed 2 of 64 treatment patients with severe pain. The output used to answer parts (a) and (c) of this problem was not affected.

(c) Since the response is ordered, we can consider a proportional odds model. However, the test for the POM has $X^2 = 21.3978$ with a P-value= 0.0110. This indicates that the POM model may not fit the data well.