## Econ 4130 2017H

## Exercises for the seminar - week 47

The exercises are based on the postponed exam 2010 H that is reproduced below for convenience (slightly edited).

## Problem 1

Introduction. When counting the number of members in groups such as the number of bacteria per colony, the number of people per household, or the number of animals per litter, every single count must necessarily be larger or equal to 1 . This excludes, e.g., the poisson distribution as a model for such counts. On the other hand the so-called logarithmic series distribution often proves useful:

A discrete random variable (rv), $Y$, taking values in $\{1,2,3, \ldots\}$, is said to be logarithmic series distributed if the probability mass function (pmf) is

$$
\begin{equation*}
P(Y=y)=f(y ; \theta)=-\frac{1}{\ln (1-\theta)} \cdot \frac{\theta^{y}}{y}, \quad y=1,2,3, \ldots \tag{1}
\end{equation*}
$$

where $\theta$ is a parameter such that $0<\theta<1$.
A. Calculate $P(Y=1)$ and $P(Y \geq 2)$ when $\theta=0.5$.
B. Show that the expected value of $Y$ is
$E(Y)=c \frac{\theta}{1-\theta} \quad$ where $\quad c=-\frac{1}{\ln (1-\theta)}$
[Hint: You are reminded of the sum of a geometric series

$$
\left.\sum_{i=0}^{\infty} a^{i}=1+a+a^{2}+a^{3}+\cdots=\frac{1}{1-a} \text { which is valid if }|a|<1\right]
$$

C. (i) Show that the moment generating function (mgf) for $Y$ is given by
$M(t)=-c \ln \left(1-\theta e^{t}\right)$, where $c$ is as given in section B. Explain why $M(t)$ is well defined in an open interval around 0 .
[Hint: You may need the following result (which you do not need to prove) from the theory of series:

$$
\left.\sum_{i=1}^{\infty} \frac{a^{i}}{i}=a+\frac{a^{2}}{2}+\frac{a^{3}}{3}+\cdots=-\ln (1-a) \text { whenever }|a|<1\right]
$$

(ii) Find $\operatorname{Var}(Y)$ as a function of $\theta$.
D. The data in table $1^{1}$ are the result of investigating the number of bacteria in each of 675 colonies of a certain type of soil bacteria. For example, the table shows that 146 of the colonies consisted of 2 bacteria, and 359 colonies had one bacterium only.

Table 1 Frequency table of the size of colonies

| Bacteria per colony $(j)$ | 1 | 2 | 3 | 4 | 5 | 6 | 7 | Sum |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Number of colonies observed <br> $f_{j}$ | 359 | 146 | 57 | 41 | 26 | 17 | 29 | 675 |

To ease the calculations we give: $\sum_{j=1}^{7} j \cdot f_{j}=1421$
Let $Y_{i}$ denote the number of bacteria in colony $i$. Assume that $Y_{1}, Y_{2}, \ldots, Y_{n}$ (where $n=675$ ) are independent and identically distributed (iid) where $Y_{i}$ is logarithmic series distributed with unknown parameter $\theta$.
(i) Show that the maximum likelihood estimator (mle) of $\theta, \hat{\theta}$, satisfies the equation

$$
\begin{equation*}
\bar{Y}=\frac{\hat{\theta}}{-(1-\hat{\theta}) \ln (1-\hat{\theta})}, \quad \text { where } \quad \bar{Y}=\frac{1}{n} \sum_{i=1}^{n} Y_{i} \tag{2}
\end{equation*}
$$

[^0][Hint: You can skip proving that the solution of (2) actually maximizes the log likelihood since the second derivative of the log likelihood is slightly complicated here. ]
(ii) Explain why the moment method estimator (mme) and the mle are the same in this case.
E. (i) Introduce the function $g(t)$ defined by
$$
g(t)=\frac{t}{-(1-t) \ln (1-t)} \text { for } 0<t<1
$$

In figure 1 we have plotted the function, $g(t)$, for $0.05<t<0.9$

Figure 1


Use the plot to obtain an approximate (rough) value for the mle estimate of $\theta$ based on the data at hand.
(ii) We are also interested in the mean colony size in the population, i.e., $E(Y)$. Explain why the maximum likelihood estimator for $E(Y)$, based on the proposed model, is simply $\hat{E}(Y)=\bar{Y}$. Present your estimate, based on the data in table 1 , of the mean number of bacteria per colony for this population.

## Problem 2

There is a complication with the data in table 1 in Problem 1. At the time when the data were collected (before 1953) it turned out difficult to count the bacteria in colonies with more than 6 bacteria. Hence all the 29 colonies registered as having 7 bacteria in table 1 should, more correctly, be registered as having 7 or more bacteria. The correct heading in the table should have been $\geq 7$ instead of only 7 . In this problem we will try to accommodate this complication.
A. The complication described above implies that the observed values of $\bar{Y}$ and the mle, $\hat{\theta}$, found in Problem 1 are not entirely correct. Are the true values of $\bar{Y}$ and $\hat{\theta}$ larger or smaller than the values found in Problem 1 when replacing the count 7 by $\geq 7$ ? Give a reason for your answer.
[Hint: You can take for granted that the function $g(t)$ is strictly increasing for $0<t<1$. Confer also figure 1 in Problem 1.]
B. Introduction. As a result of replacing the count 7 by $\geq 7$ the true value of $\bar{Y}$ cannot be calculated. But we can still estimate $E(Y)$ credibly if our chosen model for the distribution of $Y$ is realistic. A possible test of the realism of our model is the well-known Pearson $\chi^{2}$ - test. The $H_{0}$ hypothesis states that the model in Problem 1 is true (i.e., $Y_{1}, Y_{2}, \ldots, Y_{n}$ (where $n=675$ ) are iid where $Y_{i}$ is logarithmic series distributed with unknown parameter $\theta$ ). Under $H_{0}$ the frequencies in table 1 are then multinomially distributed with 7 categories, $\{1\},\{2\}, \ldots,\{6\},\{\geq 7\}$ and respective probabilities, $p_{j}(\theta)$ for $j=1,2, \ldots, 7$, where

$$
p_{j}(\theta)=\left\{\begin{array}{cl}
P(Y=j)=-\frac{1}{\ln (1-\theta)} \cdot \frac{\theta^{j}}{j} & \text { for } j=1,2, \ldots, 6 \\
P(Y \geq 7)=1-p_{1}(\theta)-p_{2}(\theta)-\cdots-p_{6}(\theta) & \text { for } j=7
\end{array}\right.
$$

Based on this multinomial specification we can determine a corrected mle, the observed value of which ${ }^{2}$ turns out to be $\hat{\theta}_{\text {obs }}=0.7569$ (you do not need to prove this).
(Note that the index obs stands for the observed value of the random variable in focus.)

Question. Perform the $\chi^{2}$-test and formulate a conclusion based on $10 \%$ level of significance. Some of the calculations necessary have been done in table 2. Complete the table by filling in the cells with question mark.

Table 2 Partial table of quantities underlying the Pearson $\chi^{2}$-test based on the corrected mle $\hat{\theta}_{\text {obs }}=0.7569^{3}$

| Category <br> $j$ <br> $j$ | Observed <br> frequency <br> $\left(O_{j}\right)$ | Mle under <br> $H_{0}$ <br> $\left(p_{j}(\hat{\theta})\right)$ | Estimated <br> frequency <br> under $H_{0}$ <br> $\left(E_{j}\right)$ | $\left(O_{j}-E_{j}\right)^{2}$ <br> $E_{j}$ <br> 1$\quad 359$ |
| :---: | :---: | :---: | :---: | :---: |
| 0.535 | 361.23 | 0.01 |  |  |
| 2 | 146 | 0.203 | 136.71 | 0.63 |
| 3 | 57 | 0.102 | 68.99 | 2.08 |
| 4 | 41 | 0.058 | 39.17 | 0.09 |
| 5 | 26 | 0.035 | 23.72 | 0.22 |
| 6 | 17 | $?$ | $?$ | $?$ |
| 7 | 29 | $?$ | $?$ | $?$ |
|  |  |  |  |  |
| Sum | 675 | 1 | 675 | $?$ |

C. (i) Explain why the likelihood function for $\theta$ used in Problem 1 cannot be used for the corrected data where category 7 is replaced by $\geq 7$.
(ii) Assuming $H_{0}$ in section $\mathbf{B}$ to be true, set up an expression for the log likelihood function for $\theta$ based on the derived multinomial model in section $\mathbf{B}$.

[^1]D. (i) Maximizing the log likelihood based on the given data, which requires numerical iterations, gives the maximum likelihood estimate, $\hat{\theta}_{\text {obs }}=0.7569$ (which you don't have to show here). Use this to compute a corrected estimate of the mean population colony size, $E(Y)$ from Problem 1. Compare with the corresponding estimate in Problem 1.
(ii) It turns out (you don't need to show this) that the Fisher information for one observation (trial) (out of $n=675$ trials ) in this (multinomial) model is $I(\theta)=\sum_{j=1}^{7} \frac{\left(p_{j}^{\prime}(\theta)\right)^{2}}{p_{j}(\theta)}$, where $p_{j}^{\prime}(\theta)$ is the derivative with respect to $\theta$ of $p_{j}(\theta)$. It turns out that the estimated value of $I(\theta)$ is
$$
I\left(\hat{\theta}_{o b s}\right)=I(0.7569)=6.6076
$$

Use this information to calculate an approximate $95 \%$ confidence interval (CI) for $\theta$ and justify the interval from general maximum likelihood theory and Slutsky's lemma.
(iii) Determine approximate $95 \%$ confidence limits for the population mean $E(Y)$ based on figure 1 in Problem 1.


[^0]:    ${ }^{1}$ Source: C.A.Bliss and R.A.Fisher, "Fitting the Negative Binomial Distribution to Biological Data", Biometrics 9 (1953): 176-200.

[^1]:    ${ }^{2}$ Determined by numerical iterative methods.
    ${ }^{3}$ The index obs stands for the observed value of the random variable in focus.

