STA310 HW3

Olivia Fu

2025-02-07

```
if (!requireNamespace("ggplot2", quietly = TRUE)) {
  install.packages("ggplot2")
}
library(ggplot2)
```

Exercise 1

The probability mass function of a Poisson random variable is given by:

$$P(X = x) = \frac{\lambda^x e^{-\lambda}}{x!}, \quad x = 0, 1, 2, \dots$$

The likelihood function is:

$$\begin{split} L(\lambda) &= \prod_{i=1}^n P(X_i = x_i) \\ &= \prod_{i=1}^n \frac{\lambda^{x_i} e^{-\lambda}}{x_i!} \\ &= \lambda^{\sum_{i=1}^n x_i} \cdot e^{-n\lambda} \cdot \prod_{i=1}^n \frac{1}{x_i!} \end{split}$$

Exercise 2

The log-likelihood function is:

$$\begin{split} \ell(\lambda) &= \log L(\lambda) \\ &= \log \left(\lambda^{\sum_{i=1}^n x_i} \cdot e^{-n\lambda} \cdot \prod_{i=1}^n \frac{1}{x_i!} \right) \\ &= \log(\lambda^{\sum_{i=1}^n x_i}) + \log(e^{-n\lambda}) + \log(\prod_{i=1}^n \frac{1}{x_i!}) \\ &= \sum_{i=1}^n x_i \cdot \log \lambda - n\lambda + \sum_{i=1}^n \log \frac{1}{x_i!} \end{split}$$

The first derivative of the log-likelihood function is:

$$\frac{\partial \ell(\lambda)}{\partial \lambda} = \frac{\sum_{i=1}^{n} x_i}{\lambda} - n$$

We then set the derivative equal to zero to find the MLE:

$$\frac{\sum_{i=1}^{n} x_i}{\hat{\lambda}} - n = 0$$

$$\hat{\lambda} = \frac{\sum_{i=1}^{n} x_i}{n}$$

The second derivative of the log-likelihood function is:

$$\frac{\partial^2 \ell(\lambda)}{\partial \lambda^2} = -\frac{\sum_{i=1}^n x_i}{\lambda^2}$$

Since $\lambda > 0$ and $x_i \ge 0$, the second derivative is negative (exclude the trivial case where all $x_i = 0$), confirming that the critical point corresponds to a maximum.

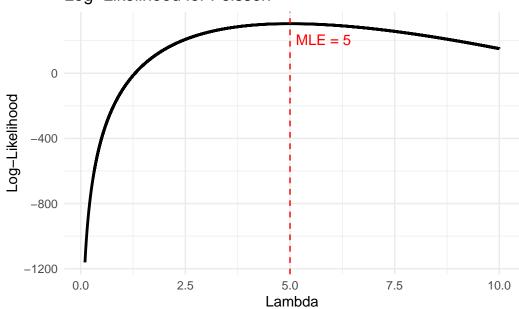
Exercise 3

```
# Given data
n <- 100
sum_x <- 500
# Log-likelihood function for Poisson distribution
log_likelihood_function <- function(lambda) {</pre>
  if (lambda <= 0) {
   return(-Inf)
  } else {
    # The constant term can be ignored as
    # it does not affect the maximization of the log-likelihood
    return(sum_x * log(lambda) - n * lambda)
  }
}
# Set up a sequence of lambda values
# Start from 0.1 to avoid lambda = 0 resulting in -Inf log-likelihood
lambda_values <- seq(0.1, 10, length.out = 100000)
# Calculate the log-likelihood for each lambda
log_likelihood_values <- sapply(lambda_values, log_likelihood_function)</pre>
# Find the lambda that maximizes the log-likelihood
mle_lambda <- lambda_values[which.max(log_likelihood_values)]</pre>
# Print the MLE for lambda
cat("MLE for lambda:", round(mle_lambda, 3), "\n")
```

MLE for lambda: 5

```
x = mle_lambda + 0.8,
y = max(log_likelihood_values) - 100,
label = paste("MLE =", round(mle_lambda, 3)),
color = "red") +
labs(title = "Log-Likelihood for Poisson",
x = "Lambda",
y = "Log-Likelihood") +
theme_minimal()
```

Log-Likelihood for Poisson



Therefore, the approximate MLE is 5, which matches the formula we derived in Exercise 2.

$$\hat{\lambda} = \frac{\sum_{i=1}^{n} x_i}{n} = \frac{500}{100} = 5$$

Exercise 4

(a)

	First five	Likelihood (no hot	
Game	shots	hand)	Likelihood (hot hand)
1	BMMBB	$p_B^3(1-p_B)^2$	$(p_B)(1-p_{B B})(1-p_B)(p_B)(p_{B B}) =$
2	MBMBM	$p_B^2(1-p_B)^3$	$ (p_B)^2 (1 - p_{B B})(1 - p_B)(p_{B B}) $ $ (1 - p_B)(p_B)(1 - p_{B B})(p_B)(1 - p_{B B}) = $
3	MMBBB	$p_{B}^{3}(1-p_{B})^{2}$	$ (p_B)^2 (1 - p_{B B})^2 (1 - p_B) $ $ (1 - p_B)(1 - p_B)(p_B)(p_{B B})(p_{B B}) = $
4	BMMMB	$p_B^2(1-p_B)^3$	$ (p_B)(1-p_B)^2(p_{B B})^2 (p_B)(1-p_{B B})(1-p_B)(1-p_B)(p_B) = $
5	MMMMM	$(1-p_B)^5$	$ (p_B)^2 (1 - p_{B B})(1 - p_B)^2 $ $ (1 - p_B)(1 - p_B)(1 - p_B)(1 - p_B)(1 - p_B) = $
			$(1 - p_B)^5$
Total		$p_B^{10}(1-p_B)^{15} \\$	$(p_B)^7 (1-p_B)^{11} (p_{B B})^3 (1-p_{B B})^4$

(b)

```
likelihood <- function(pb){
  pb^(10) * (1 - pb)^(15)
}
likelihood(0.4)</pre>
```

[1] 4.930247e-08

```
likelihood(0.3)
```

[1] 2.803388e-08

When we substitute 0.4 and 0.3 for p_B in the likelihood function, we find that $p_B = 0.4$ produces a higher likelihood than $p_B = 0.3$ (4.93 × 10⁻⁸ vs. 2.80 × 10⁻⁸). This means that the data is more consistent with $p_B = 0.4$.

Furthermore, intuitively, if the data shows that the player made 10 baskets out of 25 attempts, it makes sense to estimate the probability of making a basket close to 0.4 (10/25), rather than lower values like 0.3, which are less aligned with the observed data.

(c)

MLE for No Hot Hand Model

The likelihood and log-likelihood functions are:

$$\begin{split} Lik(p_B) &= p_B^{10} (1-p_B)^{15} \\ \log(Lik(p_B)) &= \log(p_B^{10}) + \log((1-p_B)^{15}) \\ &= 10 \log(p_B) + 15 \log(1-p_B) \end{split}$$

We then take the first derivative of the log-likelihood function and set it to zero:

$$\begin{split} \frac{d}{dp_B}log(Lik(p_B)) &= \frac{10}{p_B} - \frac{15}{1 - p_B} = 0 \\ &\Rightarrow \frac{10}{p_B} = \frac{15}{1 - p_B} \\ &\Rightarrow 10(1 - p_B) = 15p_B \\ &\Rightarrow 25p_B = 10 \\ &\hat{p_B} = \frac{10}{25} = 0.4 \end{split}$$

MLE for Hot Hand Model

The likelihood and log-likelihood functions are:

$$\begin{split} Lik(p_B, p_{B|B}) &= (p_B)^7 (1 - p_B)^{11} (p_{B|B})^3 (1 - p_{B|B})^4 \\ \log(Lik(p_B, p_{B|B})) &= \log((p_B)^7) + \log((1 - p_B)^{11}) + \log((p_{B|B})^3) + \log((1 - p_{B|B})^4) \\ &= 7 \log((p_B)) + 11 \log((1 - p_B)) + 3 \log((p_{B|B})) + 4 \log((1 - p_{B|B})) \end{split}$$

We then take the first derivative of the log-likelihood function (with respect to p_B and $p_{B|B}$ separately) and set them to zero:

$$\begin{split} \frac{d}{dp_B}log(Lik(p_B,p_{B|B})) &= \frac{7}{p_B} - \frac{11}{1-p_B} = 0 \\ &\Rightarrow \frac{7}{p_B} = \frac{11}{1-p_B} \\ &\Rightarrow 7(1-p_B) = 11p_B \\ &\Rightarrow 18p_B = 7 \\ &\hat{p_B} = \frac{7}{18} = 0.3889 \end{split}$$

$$\begin{split} \frac{d}{dp_{B|B}}log(Lik(p_B,p_{B|B})) &= \frac{3}{p_{B|B}} - \frac{4}{1 - p_{B|B}} = 0 \\ &\Rightarrow \frac{3}{p_{B|B}} = \frac{4}{1 - p_{B|B}} \\ &\Rightarrow 3(1 - p_{B|B}) = 4p_{B|B} \\ &\Rightarrow 7p_{B|B} = 3 \\ p_{B|B}^{\hat{}} &= \frac{3}{7} = 0.4286 \end{split}$$

Likelihood Ratio Test (LRT)

No Hot Hand Model: p_B

Hot Hand Model: $p_B, p_{B|B}$

Hypothesis:

$$H_0: p_B = p_{B|B}$$

$$H_a: p_B \neq p_{B|B}$$

Firstly, we need to plug the MLEs into the log-likelihood function for each model to get the maximum value of the log-likelihood for each model.

```
loglik1 <- function(pb){
  log(pb^(10) * (1 - pb)^(15))
}
loglik1(10/25)</pre>
```

[1] -16.82529

```
loglik2 <- function(pb, pbb) {
  log(pb^7 * (1-pb)^(11) * pbb^3 * (1-pbb)^4)
}
loglik2(7/18, 3/7)</pre>
```

[1] -16.80883

Then, we use the Likelihood Ratio Test to determine if the difference is statistically significant

```
LRT <- 2 * (loglik2(7/18, 3/7) - loglik1(10/25))
LRT
```

[1] 0.03292469

Therefore, LRT = 0.03292.

The test statistic follows a χ^2 distribution with 1 degrees of freedom (the difference in the number of parameters between the two models). Therefore, the p-value is $P(\chi^2 > LRT)$.

```
pchisq(LRT, 1, lower.tail = FALSE)
```

[1] 0.8560131

Therefore, p-value = 0.8560.

The p-value is very large (0.8560 > 0.05), so we fail to reject H_0 . We don't have convincing evidence that the hot hand model (conditional model) is an improvement over the no hot hand model (unconditional model). Thus, there is no significant evidence that the hot hand exists.