

# Generalized Linear Regression Homework 1

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## Problem 1

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Let the genotype frequencies be in equilibrium with probabilities:

Haptoglobin Type	Expression	Count
Hpl-1	$(1 - \theta)^2$	10
Hpl-2	$2\theta(1 - \theta)$	68
Hpl-3	$\theta^2$	112
<b>Total</b>		<b>190</b>

## Part A

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The likelihood function for the observed data can be written as,

$$L(\theta) = \binom{n}{n_1, n_2, n_3} (1 - \theta)^{2n_1} [2\theta(1 - \theta)]^{n_2} \theta^{2n_3}$$

Next, we can take the log-likelihood,

$$\ell(\theta) = n_1 \log[(1 - \theta)^2] + n_2 \log[2\theta(1 - \theta)] + n_3 \log[\theta^2]$$

Take derivative and solve for  $\theta$ ,

$$\frac{d\ell(\theta)}{d\theta} = \frac{-(2n_1 + n_2)}{1 - \theta} + \frac{n_2 + 2n_3}{\theta} = 0,$$

$$\hat{\theta} = \frac{n_2 + 2n_3}{2(n_1 + n_2 + n_3)} = 0.7684211.$$

```
# Parameters
n1 <- 10; n2 <- 68; n3 <- 112; n_total <- 190
# MLE estimate
theta_mle <- (n2 + 2 * n3) / (2 * n_total)
theta_mle
```

```
[1] 0.7684211
```



## Problem 1

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### Part B

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The Fisher information  $I(\theta)$  was solved algebraically below,

$$I(\theta) = \frac{n}{\theta(1-\theta)}$$

Thus, the asymptotic variance can be written and computed as follows,

$$\text{Var}(\hat{\theta}) = \frac{1}{I(\theta)} = \frac{\theta(1-\theta)}{n}$$

```
# Asymptotic variance
asyp_var <- theta_mle * (1 - theta_mle) / n_total
asyp_var
```

```
[1] 0.0009365797
```

Therefore, the asymptotic variance of  $\hat{\theta}$  is  $9.3657968 \times 10^{-4}$ .



## Problem 1

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### Part C

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We can use the following Wald's test statistic,

$$Z = \frac{\hat{\theta} - 0.5}{\sqrt{\text{Var}(\hat{\theta})}}$$

```
# Test statistic
z_stat <- (theta_mle - 0.5) / sqrt(asymp_var)
z_stat
```

```
[1] 8.770901
```

Using the standard normal distribution as our reference distribution,

```
p_val <- 2 * pnorm(abs(z_stat), lower.tail = FALSE)
p_val
```

```
[1] 1.772422e-18
```

Hence, the Wald  $Z$ -statistic is 8.770901, with a p-value of  $1.7724219 \times 10^{-18}$ . Thus, we reject  $H_0$ .



## Problem 2

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No, the least squares estimate would not be the same. Since regression on the mean responses of duplicates,  $\bar{Y}_i$ , removes **half of the variability** associated with each  $Y_{i,1}$  and  $Y_{i,2}$ . Even though the estimates of  $\beta_0$  and  $\beta_1$  remain unbiased, they would have a smaller variance.

Given that the two observations for each  $x_i$  are independent, the error variance can be estimated as,

$$\hat{\sigma}^2 = \frac{1}{2} \sum (Y_{i,1} - \bar{Y}_i)^2 + (Y_{i,2} - \bar{Y}_i)^2$$

Since each observation has half the variability, the estimate for the error variance remains unchanged.

## Illustration

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```
# Assume Y_i1 and Y_i2 are known
Y_i1 <- c(1, 2, 3); Y_i2 <- c(1.1, 1.9, 2.8); Y_bar <- (Y_i1 + Y_i2) / 2
# Error variance estimate
sigma_sq <- sum((Y_i1 - Y_bar)^2 + (Y_i2 - Y_bar)^2) / (2 * length(Y_i1))
sigma_sq
```

```
[1] 0.005
```

Thus,  $\hat{\sigma}^2$  is 0.005 from this specific example



## Problem 3

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Given a simple linear regression model,

$$y_i = \beta_0 + \beta_1 x_i + \epsilon_i, \quad \epsilon_i \sim N(0, \sigma^2)$$

The likelihood function for the model is based on the normality assumption of  $\epsilon_i$ . The log-likelihood function is calculated to be,

$$\ell(\beta_0, \beta_1, \sigma^2) = -\frac{n}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \sum_{i=1}^n (y_i - \beta_0 - \beta_1 x_i)^2$$

Using this, we can directly derive the MLEs for  $\beta_0$  and  $\beta_1$  by taking their respective derivatives,

### Deriving $\hat{\beta}_1$

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To find the MLE of  $\beta_1$ , we first differentiate the log-likelihood with respect to  $\beta_1$  and set it equal to zero,

$$\frac{\partial \ell}{\partial \beta_1} = -\frac{1}{\sigma^2} \sum_{i=1}^n x_i (y_i - \beta_0 - \beta_1 x_i) = 0$$

Rearranging,

$$\sum_{i=1}^n x_i y_i = \beta_0 \sum_{i=1}^n x_i + \beta_1 \sum_{i=1}^n x_i^2$$

Define  $\bar{x} = \frac{1}{n} \sum_{i=1}^n x_i$  and  $\bar{y} = \frac{1}{n} \sum_{i=1}^n y_i$ , and substitute,

$$\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y}) = \beta_1 \sum_{i=1}^n (x_i - \bar{x})^2$$

Solving for  $\beta_1$ , we get its MLE,

$$\hat{\beta}_1 = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\sum_{i=1}^n (x_i - \bar{x})^2}$$



## Deriving $\hat{\beta}_0$

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Do the same steps as above, but for  $\hat{\beta}_0$

$$\frac{\partial \ell}{\partial \beta_0} = -\frac{1}{\sigma^2} \sum_{i=1}^n (y_i - \beta_0 - \beta_1 x_i) = 0$$

Rearranging,

$$\sum_{i=1}^n y_i = n\beta_0 + \beta_1 \sum_{i=1}^n x_i$$

Simplifying,

$$\bar{y} = \beta_0 + \beta_1 \bar{x}$$

We get,

$$\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 \bar{x}$$

Thus, we have derived the MLEs for both  $\beta_0$  and  $\beta_1$ .

## Minimizing $\text{Var}(\hat{\beta}_1)$ ,

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In order to minimize  $\text{Var}(\hat{\beta}_1)$ , we need to maximize  $\sum (x_i - \bar{x})^2$ . Our best choice for the  $x_i$  values is to distribute them symmetrically and evenly across the interval  $[-1, 1]$ . We can achieve this by setting the  $x_i$  values at the endpoints,  $x_i \in \{-1, 1\}$ , or placing them symmetrically within the bounds of this interval.