# Assignment Part III

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set.seed(980824)

# Summary

#### Task 1

#### Approach:

To calculate the AIC, the formula

$$2 * k - 2 * l(\hat{\theta_{ml}})$$

was applied where k is the number of parameters in the theta. To calculate the leave-one-out cross-validation, the formula

$$l(\hat{\theta_{ml}}) + \sum_{i=1}^{n} ((S_i(\theta_i)^T) * (\hat{\theta_i} - \hat{\theta_{ml}}))$$

was applied where  $\hat{\theta_i}$  is  $\hat{\theta_{ml}}(X_{-i})$ ,  $(X_{-i})$  is one observation  $(X_i)$  is left out and n is the total number of observations. These values were then compared with the AIC from R in the provided summary.

#### Code:

```
# ---- Task_1 ----
# Compute AIC = 2k - 2l(theta_ml)
n \leftarrow nrow(X)
theta0 = rep(0, ncol(X))
k <- length(theta0)</pre>
theta_estimate <- NR(theta0, 3, y, X)</pre>
log_likellihood <- 1(theta_estimate, y, X)</pre>
aic_computed <- 2*k - 2*log_likellihood
#AIC output from R summary
r_summary_aic <- summary(modell)$aic</pre>
 \# \ \textit{Compute} \ n*k\_cv = l(\textit{theta\_ml}) + \textit{sum}((S\_i(\textit{theta\_i}))T*(\textit{theta\_i} - \textit{theta\_ml})) 
# Here, theta_i = theta_ml(X_-i)
nk_cv <- log_likellihood
for(i in 1:n) {
  X_minus_i <- X[-i, , drop=FALSE]</pre>
  y_minus_i <- y[-i, , drop=FALSE]</pre>
  theta_i <- NR(theta0, 3, y_minus_i, X_minus_i)</pre>
  score <- S(theta_i, y_minus_i, X_minus_i)</pre>
```

```
subset_sum <- t(score) %*% (theta_i - theta_estimate)
nk_cv <- nk_cv + subset_sum
}

# Creating a comparison data frame
comparison_aic_values <- data.frame(
    "AIC_R_model" = r_summary_aic,
    "AIC_computed" = aic_computed,
    "NK_CV_computed" = -2*nk_cv
)</pre>
```

## Output:

```
comparison_aic_values
```

```
## AIC_R_model AIC_computed NK_CV_computed ## 1 1302.397 1302.397 1294.397
```

### Task 2

Approach:

Code:

```
# ---- Task_2 ----
```

Output:

Observation:

Task 3

Approach:

Code:

```
# ---- Task_3 ----
```

Output:

Observation: