In [1]:
```r
# Re-running the code for Glm after Bootstrapping
# NOTE : Star rating part of the program not required . Star is displayed automa

# Load your data
data <- read.csv("Complete_Data_Modified.csv")
#data <- data2
# Define your variables
independent_variables <- c('BFI..E.', 'BFI..A.', 'BFI..C.', 'BFI..N.', 'BFI..O.'
mediator_variable <- 'PAQ'
dependent_variables <- c('Panas..', 'Panas...1', 'CBCL')

# Function to calculate star ratings
get_star_rating <- function(p_value) {
  if (p_value < 0.001) return("***")
  else if (p_value < 0.01) return("**")
  else if (p_value < 0.05) return("*")
  else return("check")
}

# Fit separate Poisson GLM models for each dependent variable
for (dependent_variable in dependent_variables) {
  formula <- paste(dependent_variable, "~ . +", mediator_variable)
  poisson_model <- glm(formula, data = data, family = poisson(link = "log"))
  print(poisson_model)
  # Summarize the model
  cat("Summary for", dependent_variable, ":\n")
  model_summary <- summary(poisson_model)
  print(model_summary)


  # Get and print star ratings for p-values
  p_values <- model_summary$coefficients[, "Pr(>|z|)"]
  star_ratings <- sapply(p_values, get_star_rating)
  star_ratings_df <- data.frame(Coefficient = rownames(model_summary$coefficient
  print(star_ratings_df)
  print("=====================================================================
}
```

```
Call:  glm(formula = formula, family = poisson(link = "log"), data = data)

Coefficients:
  (Intercept)        Panas...1         BFI..E.          BFI..A.          BFI..C.
    0.7700304        0.0135235        0.0124698        0.1079799       -0.0856901
       BFI..N.          BFI..O.       EI..Self.A.      EI..Self.M.     EI..Social.A.
   -0.0047829        0.1434257        0.0004412        0.0040463       -0.0043248
        EI..RM.            Total             PAQ             CBCL
   -0.0037535        0.0097494       -0.0831965       -0.1125351

Degrees of Freedom: 205 Total (i.e. Null);  192 Residual
Null Deviance:        43.96
Residual Deviance: 37.48          AIC: 552.2
Summary for Panas.. :

Call:
glm(formula = formula, family = poisson(link = "log"), data = data)

Coefficients:
                Estimate Std. Error z value Pr(>|z|)
(Intercept)    0.7700304  1.2265366   0.628    0.530
Panas...1      0.0135235  0.1386784   0.098    0.922
BFI..E.        0.0124698  0.1284255   0.097    0.923
BFI..A.        0.1079799  0.1262575   0.855    0.392
BFI..C.       -0.0856901  0.1350942  -0.634    0.526
BFI..N.       -0.0047829  0.1174790  -0.041    0.968
BFI..O.        0.1434257  0.1621923   0.884    0.377
EI..Self.A.    0.0004412  0.0088393   0.050    0.960
EI..Self.M.    0.0040463  0.0075069   0.539    0.590
EI..Social.A. -0.0043248  0.0095737  -0.452    0.651
EI..RM.       -0.0037535  0.0077426  -0.485    0.628
Total          0.0097494  0.1626609   0.060    0.952
PAQ           -0.0831965  0.0799855  -1.040    0.298
CBCL          -0.1125351  0.0768828  -1.464    0.143


(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 43.958  on 205  degrees of freedom
Residual deviance: 37.477  on 192  degrees of freedom
AIC: 552.21

Number of Fisher Scoring iterations: 4


                 Coefficient Stars
(Intercept)        (Intercept) check
Panas...1            Panas...1 check
BFI..E.                BFI..E. check
BFI..A.                BFI..A. check
BFI..C.                BFI..C. check
BFI..N.                BFI..N. check
BFI..O.                BFI..O. check
EI..Self.A.        EI..Self.A. check
EI..Self.M.        EI..Self.M. check
EI..Social.A.    EI..Social.A. check
EI..RM.                EI..RM. check
Total                    Total check
PAQ                        PAQ check
CBCL                      CBCL check
[1] "==============================================================="
```

```
Call: glm(formula = formula, family = poisson(link = "log"), data = data)

Coefficients:
  (Intercept)         Panas..          BFI..E.         BFI..A.         BFI..C.
    -0.609854        0.010821        -0.089217        0.030407        0.099554
      BFI..N.          BFI..O.       EI..Self.A.     EI..Self.M.    EI..Social.A.
     0.035063        -0.008884        0.001150        0.003087        0.012130
       EI..RM.           Total              PAQ            CBCL
    -0.006606        -0.034919        -0.045686        0.039844

Degrees of Freedom: 205 Total (i.e. Null);  192 Residual
Null Deviance:        26.32
Residual Deviance: 22.95        AIC: 491.8
Summary for Panas...1 :

Call:
glm(formula = formula, family = poisson(link = "log"), data = data)

Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)  -0.609854   1.424399  -0.428    0.669
Panas..       0.010821   0.117047   0.092    0.926
BFI..E.      -0.089217   0.143465  -0.622    0.534
BFI..A.       0.030407   0.143949   0.211    0.833
BFI..C.       0.099554   0.152793   0.652    0.515
BFI..N.       0.035063   0.134671   0.260    0.795
BFI..O.      -0.008884   0.184830  -0.048    0.962
EI..Self.A.   0.001150   0.009946   0.116    0.908
EI..Self.M.   0.003087   0.008709   0.355    0.723
EI..Social.A. 0.012130   0.011133   1.090    0.276
EI..RM.      -0.006606   0.008778  -0.753    0.452
Total        -0.034919   0.184343  -0.189    0.850
PAQ          -0.045686   0.090015  -0.508    0.612
CBCL          0.039844   0.095055   0.419    0.675


(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 26.322  on 205  degrees of freedom
Residual deviance: 22.946  on 192  degrees of freedom
AIC: 491.79

Number of Fisher Scoring iterations: 4


                  Coefficient Stars
(Intercept)       (Intercept) check
Panas..               Panas.. check
BFI..E.               BFI..E. check
BFI..A.               BFI..A. check
BFI..C.               BFI..C. check
BFI..N.               BFI..N. check
BFI..O.               BFI..O. check
EI..Self.A.       EI..Self.A. check
EI..Self.M.       EI..Self.M. check
EI..Social.A. EI..Social.A. check
EI..RM.               EI..RM. check
Total                   Total check
PAQ                       PAQ check
CBCL                     CBCL check
[1] "==============================================================="
```

```
Call:  glm(formula = formula, family = poisson(link = "log"), data = data)

Coefficients:
  (Intercept)         Panas..        Panas...1         BFI..E.         BFI..A.
    1.3283255      -0.1171280        0.0528894       0.0241293      -0.0437459
        BFI..C.         BFI..N.         BFI..O.     EI..Self.A.     EI..Self.M.
    -0.0756876       0.0409897      -0.0332714      -0.0005178       0.0001310
EI..Social.A.         EI..RM.           Total             PAQ
   -0.0013461      -0.0018969        0.0351775      -0.0148301

Degrees of Freedom: 205 Total (i.e. Null);  192 Residual
Null Deviance:       50
Residual Deviance: 44.27        AIC: 639.6
Summary for CBCL :

Call:
glm(formula = formula, family = poisson(link = "log"), data = data)

Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)   1.3283255  0.9831334   1.351    0.177
Panas..      -0.1171280  0.0801852  -1.461    0.144
Panas...1     0.0528894  0.1104107   0.479    0.632
BFI..E.       0.0241293  0.1028378   0.235    0.814
BFI..A.      -0.0437459  0.1021794  -0.428    0.669
BFI..C.      -0.0756876  0.1077143  -0.703    0.482
BFI..N.       0.0409897  0.0949686   0.432    0.666
BFI..O.      -0.0332714  0.1334628  -0.249    0.803
EI..Self.A.  -0.0005178  0.0071807  -0.072    0.943
EI..Self.M.   0.0001310  0.0062098   0.021    0.983
EI..Social.A. -0.0013461  0.0076985  -0.175    0.861
EI..RM.      -0.0018969  0.0064075  -0.296    0.767
Total         0.0351775  0.1328631   0.265    0.791
PAQ          -0.0148301  0.0626697  -0.237    0.813


(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 50.005  on 205  degrees of freedom
Residual deviance: 44.267  on 192  degrees of freedom
AIC: 639.64

Number of Fisher Scoring iterations: 4

                 Coefficient Stars
(Intercept)       (Intercept) check
Panas..              Panas.. check
Panas...1          Panas...1 check
BFI..E.              BFI..E. check
BFI..A.              BFI..A. check
BFI..C.              BFI..C. check
BFI..N.              BFI..N. check
BFI..O.              BFI..O. check
EI..Self.A.      EI..Self.A. check
EI..Self.M.      EI..Self.M. check
EI..Social.A. EI..Social.A. check
EI..RM.              EI..RM. check
Total                  Total check
PAQ                      PAQ check
[1] "================================================================="
```

In [ ]:
```r
# Bootstrapping code to run for 300 iterations . Can be expanded later

# Load your data (assuming 'data' is already loaded)
# data <- read.csv("your_data.csv")
data <- read.csv("Complete_Data_Modified.csv")

# Define your variables
independent_vars <- c('BFI..E.', 'BFI..A.', 'BFI..C.', 'BFI..N.', 'BFI..O.', 'EI
dependent_vars <- c('Panas..', 'Panas...1', 'CBCL')
mediator_var <- 'PAQ'

# Set the new sample size , here taken 150
new_sample_size <- 600

# Create an empty data frame to store the bootstrapped data
data2 <- data.frame(matrix(NA, ncol = length(independent_vars) + length(dependen
colnames(data2) <- c(independent_vars, dependent_vars, mediator_var)

# Perform bootstrapping
for (i in 1:new_sample_size) {
  # Generate a random sample with replacement from the original data
  random_indices <- sample(nrow(data), replace = TRUE)
  bootstrap_sample <- data[random_indices, ]

  # Store the bootstrap sample in data2
  data2[i, ] <- bootstrap_sample
}

# Now data2 contains the bootstrapped data with the new sample size
data <- data2
# Define your variables
independent_variables <- c('BFI..E.', 'BFI..A.', 'BFI..C.', 'BFI..N.', 'BFI..O.'
mediator_variable <- 'PAQ'
dependent_variables <- c('Panas..', 'Panas...1', 'CBCL')

# Function to calculate star ratings
get_star_rating <- function(p_value) {
  if (p_value < 0.001) return("***")
  else if (p_value < 0.01) return("**")
  else if (p_value < 0.05) return("*")
  else return("check")
}

# Fit separate Poisson GLM models for each dependent variable
for (dependent_variable in dependent_variables) {
  formula <- paste(dependent_variable, "~ . +", mediator_variable)
  poisson_model <- glm(formula, data = data, family = poisson(link = "log"))
  print(poisson_model)
  # Summarize the model
  cat("Summary for", dependent_variable, ":\n")
  model_summary <- summary(poisson_model)
  print(model_summary)


  # Get and print star ratings for p-values
  p_values <- model_summary$coefficients[, "Pr(>|z|)"]
  star_ratings <- sapply(p_values, get_star_rating)
  star_ratings_df <- data.frame(Coefficient = rownames(model_summary$coefficient
  print(star_ratings_df)
```

```
  print("==============================================================
}
```

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s.. = c(1L, :
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Regression_Linear_Possion

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In [ ]: