In [1]:
```r
#Sources used ::
# 1) https://www.dataquest.io/blog/tutorial-poisson-regression-in-r/
# 2) https://timeseriesreasoning.com/contents/poisson-regression-model/
# 3) https://cran.r-project.org/web/packages/sjPlot/vignettes/tab_model_estimates.h
# 4) https://online.stat.psu.edu/stat504/lesson/9/9.2
# 5) https://www.geeksforgeeks.org/poisson-regression-in-r-programming/ (Most used

# Load your data
data <- read.csv("Pilot_modified_data_1.csv")

# 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"))

  # 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$coefficients[, "Pr(>|z|)"]
  star_ratings <- sapply(p_values, get_star_rating)
  star_ratings_df <- data.frame(Coefficient = rownames(model_summary$coefficients),
  print(star_ratings_df)
  print("================================================================")
}
```

Summary for Panas.. :

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

Deviance Residuals:
      Min   1Q   Median   3Q   Max
 -0.127254 -0.030803 -0.004841   0.030804   0.236661

Coefficients:
                 Estimate Std. Error z value Pr(>|z|)
(Intercept)   0.8376069 2.7176884   0.308   0.758
Panas...1   -0.0035951 0.2941212 -0.012   0.990
BFI..E.   -0.0490939 0.4710388 -0.104   0.917
BFI..A.   0.0071564 0.2813401   0.025   0.980
BFI..C.   0.0366116 0.4633167   0.079   0.937
BFI..N.   0.0525955 0.2210784   0.238   0.812
BFI..O.   -0.0122170 0.3629494 -0.034   0.973
EI..Self.A.   0.0008079 0.0183715   0.044   0.965
EI..Self.M.   -0.0021908 0.0143532 -0.153   0.879
EI..Social.A.   0.0023679 0.0235086   0.101   0.920
EI..RM.   -0.0001474 0.0164597 -0.009   0.993
Total   -0.0173123 0.3458867 -0.050   0.960
PAQ   0.0146337 0.2688049   0.054   0.957
CBCL   -0.1406602 0.4055748 -0.347   0.729

(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 0.41816 on 33 degrees of freedom
Residual deviance: 0.15941 on 20 degrees of freedom
AIC: 117.4

Number of Fisher Scoring iterations: 3

                    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] "=="
Summary for Panas...1 :

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

Deviance Residuals:

```
    Min      1Q   Median      3Q      Max
-0.6553  -0.3270  -0.0017   0.2423   0.5873
```

Coefficients:

	Estimate	Std. Error	z value	Pr(>|z|)
(Intercept)	0.889891	4.349013	0.205	0.838
Panas..	-0.149219	1.404358	-0.106	0.915
BFI..E.	-0.146126	0.531573	-0.275	0.783
BFI..A.	-0.033461	0.324314	-0.103	0.918
BFI..C.	0.154399	0.543706	0.284	0.776
BFI..N.	0.062337	0.295745	0.211	0.833
BFI..O.	-0.280332	0.419729	-0.668	0.504
EI..Self.A.	0.003700	0.020197	0.183	0.855
EI..Self.M.	-0.000276	0.017381	-0.016	0.987
EI..Social.A.	0.002127	0.027020	0.079	0.937
EI..RM.	-0.007422	0.017756	-0.418	0.676
Total	0.060747	0.382359	0.159	0.874
PAQ	0.082828	0.304525	0.272	0.786
CBCL	0.089480	0.656400	0.136	0.892

(Dispersion parameter for poisson family taken to be 1)

```
    Null deviance: 5.7187  on 33  degrees of freedom
Residual deviance: 3.8829  on 20  degrees of freedom
AIC: 110.93
```

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] "==="
Summary for CBCL :

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

Deviance Residuals:
```
     Min       1Q   Median       3Q      Max
-0.60355  -0.06141  0.02928  0.07397  0.28235
```

Coefficients:

	Estimate	Std. Error	z value	Pr(>|z|)
(Intercept)	2.104e+00	3.339e+00	0.630	0.529

```
Panas..          -8.260e-01  1.114e+00  -0.741     0.459
Panas...1         2.974e-02  3.034e-01   0.098     0.922
BFI..E.          -1.124e-01  4.808e-01  -0.234     0.815
BFI..A.          -4.170e-02  2.832e-01  -0.147     0.883
BFI..C.           1.188e-01  4.719e-01   0.252     0.801
BFI..N.           9.358e-02  2.512e-01   0.373     0.709
BFI..O.          -7.562e-02  3.670e-01  -0.206     0.837
EI..Self.A.       1.366e-04  1.894e-02   0.007     0.994
EI..Self.M.      -3.774e-03  1.527e-02  -0.247     0.805
EI..Social.A.     7.694e-03  2.340e-02   0.329     0.742
EI..RM.          -1.222e-05  1.706e-02  -0.001     0.999
Total            -9.119e-02  3.408e-01  -0.268     0.789
PAQ               6.153e-02  2.703e-01   0.228     0.820

(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 2.2148  on 33  degrees of freedom
Residual deviance: 1.1489  on 20  degrees of freedom
AIC: 115.56

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 [2]:
```r
# Bootstrapping code to run for 150 iterations . Can be expanded later

# Load your data (assuming 'data' is already loaded)
# data <- read.csv("your_data.csv")
data <- read.csv("Pilot_modified_data_1.csv")

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

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

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

```r
# 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
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 2 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 3 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 4 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 6 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 13 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 14 has 34 rows to replace 1 rows"
```

```
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 3 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 4 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 7 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 8 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 9 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 10 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 11 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 12 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 13 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 14 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 1 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 2 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 3 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 4 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
```

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"replacement element 5 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 6 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 7 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 8 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 9 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
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Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 11 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 12 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
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"replacement element 13 has 34 rows to replace 1 rows"
Warning message in `[<-.data.frame`(`*tmp*`, i, , value = structure(list(Panas.. =
c(2L, :
"replacement element 14 has 34 rows to replace 1 rows"

</div>

In [3]:
```r
# Re-running the code for Glm after Bootstrapping
# NOTE : Star rating part of the program not required . Star is displayed automatic

# Load your data
data <- read.csv("Pilot_modified_data_1.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"))

  # Summarize the model
  cat("Summary for", dependent_variable, ":\n")
  model_summary <- summary(poisson_model)
```

```r
  print(model_summary)


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

Summary for Panas.. :

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

Deviance Residuals:
    Min   1Q   Median   3Q   Max
 -2.1613 -0.4921   0.3011   0.4881   2.4181

Coefficients:
                 Estimate Std. Error z value Pr(>|z|)
(Intercept)   3.489093   0.342497 10.187 < 2e-16 ***
BFI..E.   0.035017   0.115094   0.304 0.760943
BFI..A.   -0.099170   0.024187 -4.100 4.13e-05 ***
BFI..C.   -0.068885   0.033460 -2.059 0.039519 *
BFI..N.   0.015480   0.023646   0.655 0.512705
BFI..O.   0.023444   0.036183   0.648 0.517028
EI..Self.A.   0.013581   0.022691   0.599 0.549492
EI..Self.M.   -0.043974   0.030118 -1.460 0.144270
EI..Social.A.   0.008153   0.001489   5.476 4.35e-08 ***
EI..RM.   -0.004767   0.001166 -4.090 4.32e-05 ***
Total   0.002949   0.002099   1.405 0.160166
Panas...1   0.130386   0.025346   5.144 2.69e-07 ***
CBCL   -0.072658   0.021319 -3.408 0.000654 ***
PAQ   0.095267   0.054744   1.740 0.081817 .

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 336.761 on 149 degrees of freedom
Residual deviance: 89.451 on 136 degrees of freedom
AIC: 1027.1

Number of Fisher Scoring iterations: 4

                  Coefficient Stars
(Intercept)   (Intercept)   ***
BFI..E.   BFI..E. check
BFI..A.   BFI..A.   ***
BFI..C.   BFI..C.   *
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.   ***
EI..RM.   EI..RM.   ***
Total   Total check
Panas...1   Panas...1   ***
CBCL   CBCL   ***
PAQ   PAQ check
[1] "==="
Summary for Panas...1 :

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

```
Deviance Residuals:
     Min        1Q     Median        3Q        Max
-0.44531   -0.15188    0.01827    0.10269    0.65584


Coefficients:
               Estimate Std. Error z value Pr(>|z|)
(Intercept)    -0.430275   1.719042  -0.250    0.802
BFI..E.        -0.277772   0.628896  -0.442    0.659
BFI..A.         0.041187   0.125283   0.329    0.742
BFI..C.        -0.027574   0.170353  -0.162    0.871
BFI..N.        -0.015721   0.123225  -0.128    0.898
BFI..O.         0.111027   0.190332   0.583    0.560
EI..Self.A.     0.040508   0.117661   0.344    0.731
EI..Self.M.    -0.098317   0.156265  -0.629    0.529
EI..Social.A.  -0.004249   0.008503  -0.500    0.617
EI..RM.         0.007271   0.005530   1.315    0.189
Total           0.017990   0.008237   2.184    0.029 *
Panas..         0.010347   0.006645   1.557    0.119
CBCL            0.096129   0.106178   0.905    0.365
PAQ            -0.274413   0.247676  -1.108    0.268
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1


(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 44.7824  on 149  degrees of freedom
Residual deviance:  6.6684  on 136  degrees of freedom
AIC: 462.96


Number of Fisher Scoring iterations: 4


               Coefficient Stars
(Intercept)      (Intercept) 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     *
Panas..              Panas.. check
CBCL                    CBCL check
PAQ                      PAQ check
[1] "======================================================================="
Summary for CBCL :


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


Deviance Residuals:
     Min        1Q     Median        3Q        Max
-0.94840   -0.25428   -0.07813    0.23530    0.75006
```

```
Coefficients:
                 Estimate Std. Error z value Pr(>|z|)
(Intercept)     -0.499935   2.142848  -0.233 0.815525
BFI..E.         -0.006020   0.723273  -0.008 0.993359
BFI..A.          0.102724   0.173604   0.592 0.554042
BFI..C.          0.024343   0.253217   0.096 0.923413
BFI..N.          0.400457   0.120142   3.333 0.000859 ***
BFI..O.         -0.243300   0.237796  -1.023 0.306237
EI..Self.A.      0.046577   0.157702   0.295 0.767727
EI..Self.M.      0.148720   0.188665   0.788 0.430534
EI..Social.A.    0.008083   0.010309   0.784 0.432981
EI..RM.         -0.004351   0.007555  -0.576 0.564649
Total           -0.006752   0.014522  -0.465 0.641970
Panas..         -0.012027   0.009195  -1.308 0.190901
Panas...1        0.220227   0.171162   1.287 0.198214
PAQ              0.218956   0.374334   0.585 0.558601
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1


(Dispersion parameter for poisson family taken to be 1)

    Null deviance: 39.484  on 149  degrees of freedom
Residual deviance: 15.773  on 136  degrees of freedom
AIC: 405.58

Number of Fisher Scoring iterations: 4


                 Coefficient Stars
(Intercept)      (Intercept) check
BFI..E.              BFI..E. check
BFI..A.              BFI..A. check
BFI..C.              BFI..C. check
BFI..N.              BFI..N.   ***
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
Panas..              Panas.. check
Panas...1          Panas...1 check
PAQ                      PAQ check
[1] "================================================================="
```

In [ ]: