

What is Zero-truncated Negative Binomial Regression, and how is it used in Mplus data analysis?

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Zero-truncated Negative Binomial Regression is a statistical technique used in Mplus data analysis to model count data with excess zeros. It is a type of regression analysis that is suitable for analyzing data with a high proportion of zero values, where the traditional Negative Binomial Regression may not be appropriate. This method takes into account the truncation of the data at zero, which means it only models the counts above zero. It is often used in social science research to analyze data such as number of absences, number of hospitalizations, or number of criminal offenses. By incorporating the excess zeros, this method provides a more accurate estimation of the relationship between the independent variables and the count outcome. It is particularly useful in analyzing data with overdispersion, where the variance is greater than the mean. In summary, Zero-truncated Negative Binomial Regression is a specialized statistical tool that allows for the analysis of count data with a high proportion of zeros, providing researchers with a more precise understanding of the relationships between variables in their data.

Zero-truncated Negative Binomial Regression | Mplus Data Analysis Examples

Version info: Code for this page was tested in Mplus version 6.12.

Zero-truncated negative binomial regression is used to model count data for which the value zero cannot occur and when there is evidence of over dispersion .

Please Note: The purpose of this page is to show how to use various data analysis commands.

It does not cover all aspects of the research process which researchers are expected to do. In particular, it does not cover data cleaning and

verification, verification of assumptions, model diagnostics and potential follow-up analyses.

Examples of zero-truncated negative binomial

Example 1.

A study of the length of hospital stay, in days, as a function of age, kind of health insurance and whether or not the patient died while in the hospital.

Length of hospital stay is recorded as a minimum of at least one day.

Example 2.

A study of the number of journal articles published by tenured faculty as a function of discipline (fine arts, science, social science, humanities, medical, etc). To get tenure faculty must publish, i.e., there are no tenured faculty with zero publications.

Example 3.

A study by the county traffic court on the number of

tickets received by teenagers as predicted by school performance, amount of driver training and gender. Only individuals who have received at least one citation are in the traffic court files.

Description of the data

Let's pursue Example 1 from above.

We have a hypothetical data file available here with 1,493 observations.

The variable describing length of hospital visit is stay. The variable age gives the age group from 1 to 9 which will be treated as interval in this example.

The variables hmo and died are binary indicator variables for HMO insured patients and patients who died while in hospital, respectively.

Let's look at the data.

Data:

File is C:ztnb.dat;

Variable:

Names are

stay age hmo died;

Missing are all (-9999) ;

Analysis:

Type = basic ;

Plot:

Type = plot1;

ESTIMATED SAMPLE STATISTICS

Means

STAY AGE HMO DIED

1 9.729 5.234 0.160 0.343

Covariances

STAY AGE HMO DIED

STAY 66.100

AGE -0.615 2.785

HMO -0.169 -0.006 0.134

DIED -0.447 0.121 0.000 0.225

Correlations

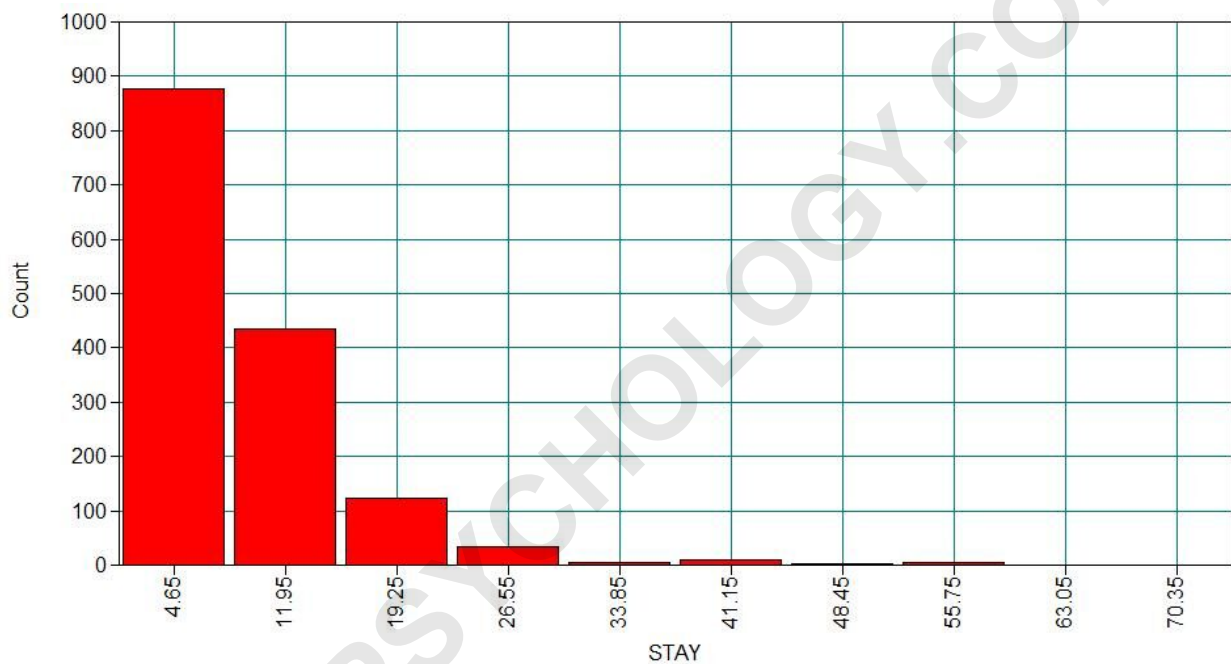
STAY AGE HMO DIED

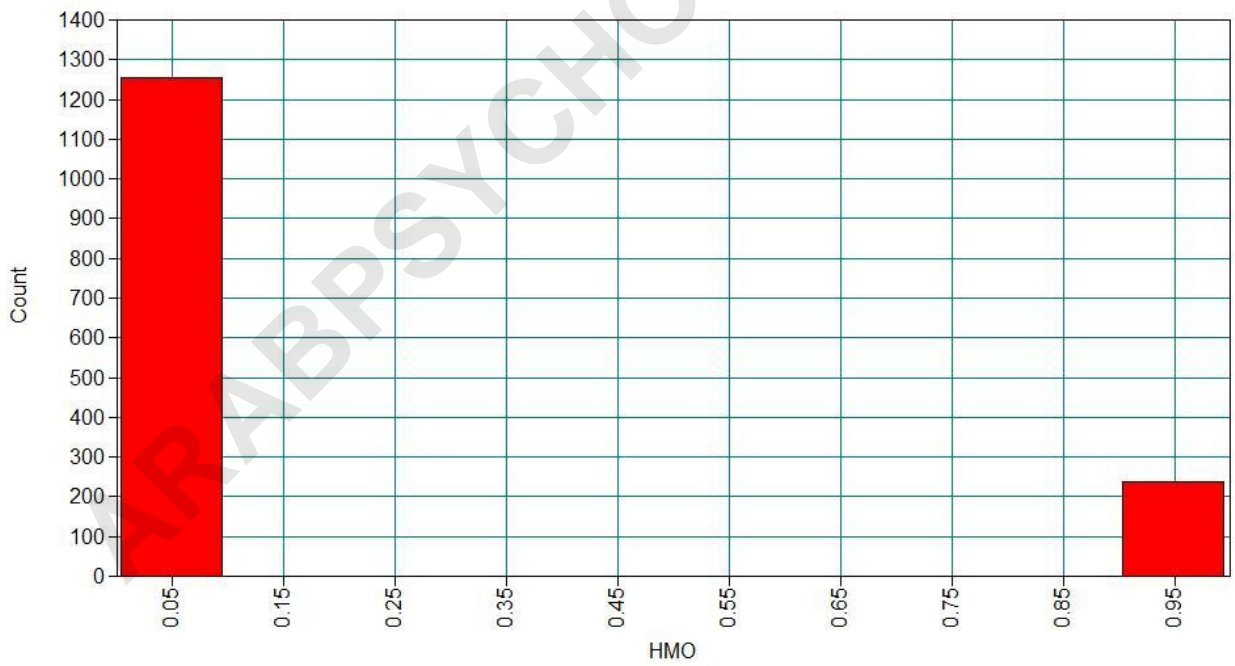
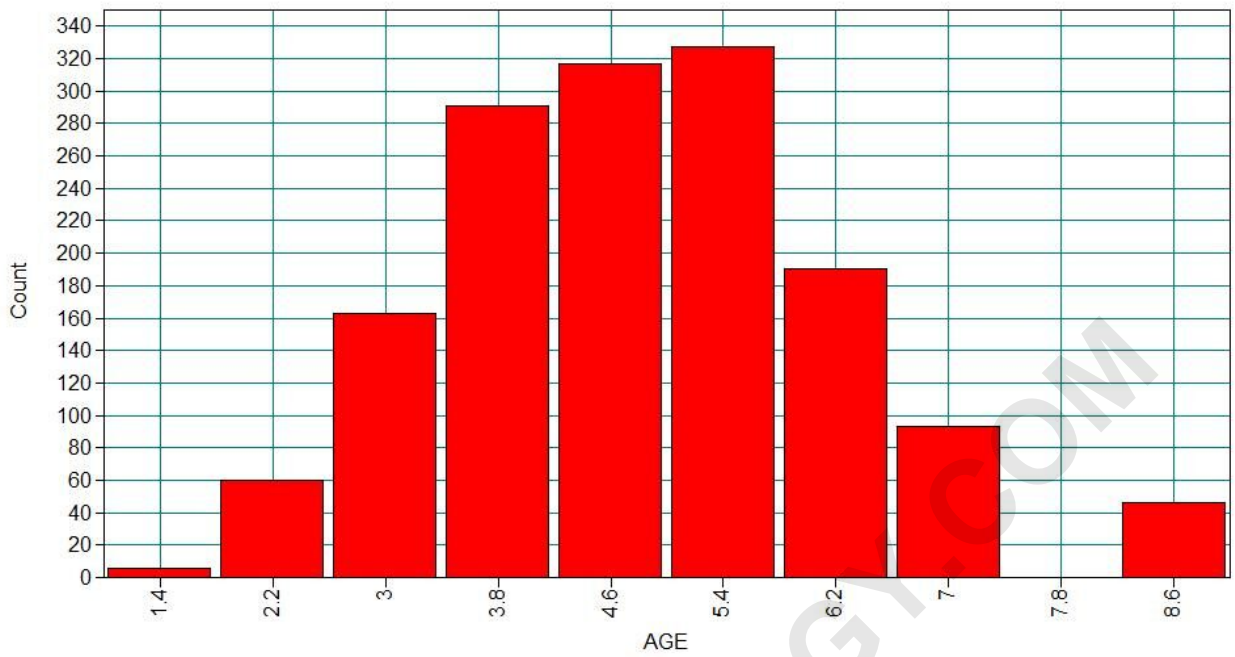
STAY 1.000

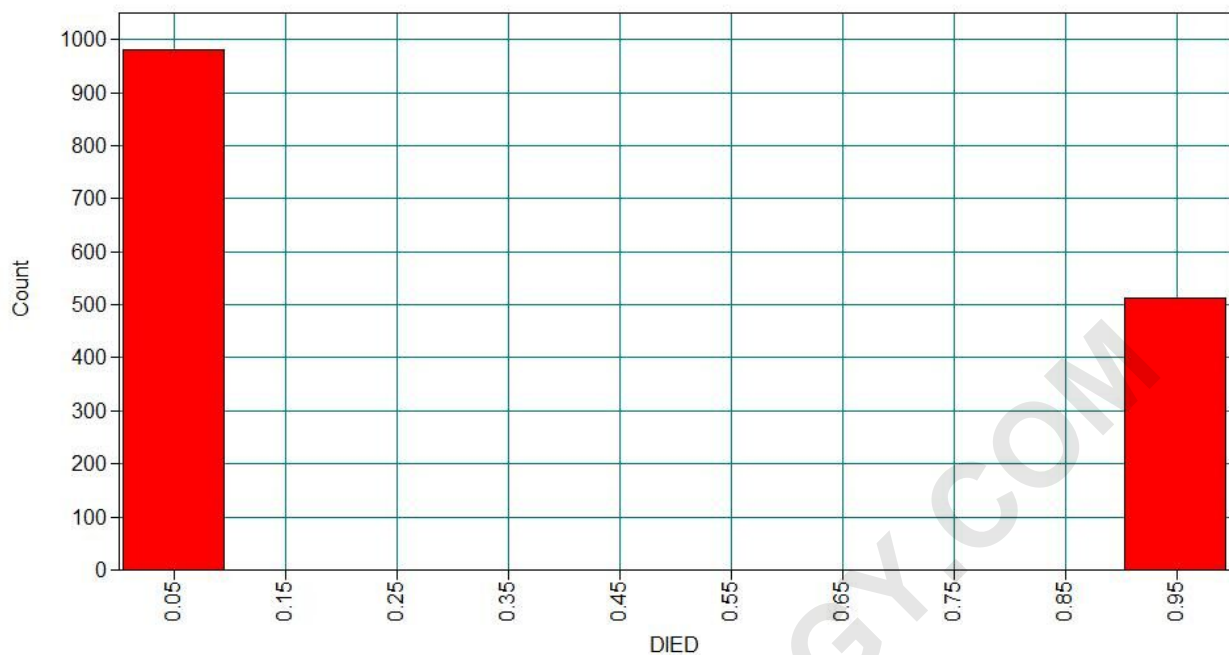
AGE -0.045 1.000

HMO -0.057 -0.010 1.000

DIED -0.116 0.152 0.000 1.000







Analysis methods you might consider

Before we show how you can analyze these data with a zero-truncated negative binomial analysis, let's consider some other methods that you might use.

Zero-truncated negative binomial regression

In the syntax below, we have indicated that stay is a count variable by using the count statement. The (nbt) option is used to indicate 2 things: that we are modeling our

count variable with a negative binomial distribution, and that we are specifying a zero-truncated model.

Without the (t) option we would be estimating a negative

binomial model without

zero-truncation. Also, we do not need a usevariables statement

because

we are using all of the variables in the data set in the current model.

We have omitted the missing statement because we have no missing data in

this data set. The default estimation method is MLR - maximum likelihood

parameter estimates with standard errors and a chi-square test statistic that

are robust to non-normality and non-independence of observations. The MLR standard errors

are computed using a sandwich estimator. This is what we generally call robust

standard errors. To get the "regular" standard errors, we use the estimator

= ml on the analysis statement. (In the next example, we

will

omit the analysis statement and obtain the robust standard errors.)

Our regression equations is specified in the model statement: we are predicting length of stay using age, hmo status and whether the patient died.

Data:

File is C:ztnb.dat ;

Variable:

Names = stay age hmo died;

Count = stay(nbt);

Model:

stay on age hmo died;

MODEL FIT INFORMATION

Number of Free Parameters 5

Loglikelihood

H0 Value -4755.280

H0 Scaling Correction Factor 1.156

for MLR

Information Criteria

Akaike (AIC) 9520.559

Bayesian (BIC) 9547.102

Sample-Size Adjusted BIC 9531.218

($n^* = (n + 2) / 24$)

MODEL RESULTS

Two-Tailed

Estimate S.E. Est./S.E. P-Value

STAY ON

AGE -0.016 0.013 -1.194 0.233

HMO -0.147 0.057 -2.571 0.010

DIED -0.218 0.053 -4.142 0.000

Intercepts

STAY 2.408 0.075 32.039 0.000

Dispersion

STAY 0.566 0.037 15.316 0.000

In the MODEL FIT INFORMATION portion of the output, you will find the log likelihood for the final model as well as a number of fit

statistics. In the **MODEL RESULTS** section of the output you will find the negative binomial regression coefficients (estimates) for each of the variables, standard errors and the ratio of the estimate to its standard error. This can be used as a Z test, where values greater than 2 are considered to be statistically significant. We see that hmo and died but not age are significant predictors of stay. Thus, for example, for patients who use HMO services compared to those who do not, the log count of days stayed is about 0.147 less.

Now let's rerun the model without the analysis: estimator = ml statement in order to obtain robust standard errors.

Data:

File is C:ztnb.dat ;

Variable:

Names = stay age hmo died;

Missing = all (-9999) ;

Count = stay(nbt);

Model:

stay on age hmo died;

Analysis:

estimator = ml;

MODEL RESULTS

Two-Tailed

Estimate S.E. Est./S.E. P-Value

STAY ON

AGE -0.016 0.013 -1.197 0.231

HMO -0.147 0.059 -2.483 0.013

DIED -0.218 0.046 -4.718 0.000

Intercepts

STAY 2.408 0.072 33.457 0.000

Dispersion

STAY 0.566 0.031 18.132 0.000

Robust standard errors tend to be larger than "regular" standard errors,

though not always as we see for the variable age. The results changed very little when using regular standard errors.

Things to consider

References

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