



## ggESDA: An R Package for exploratory symbolic data analysis using ggplot2

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### Abstract

This paper presents the **ggESDA** package, which we developed for exploratory symbolic data analysis in R. Based on **ggplot2** Wickham (2009), the **ggESDA** package which is familiar

*Keywords:* SDA, EDA, symbolic data analysis, exploratory data analysis, ggplot2 extensions, interval-valued data, R.

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## 1. Introduction: xxx(wait for edit)

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R (wait for edit) `glm()` (Chambers and Hastie 1992) in the **stats** package.

```
glm(formula, data, subset, na.action, weights, offset,  
     family = gaussian, start = NULL, control = glm.control(...),  
     model = TRUE, y = TRUE, x = FALSE, ...)
```

Type	Distribution	Method	Description
GLM	Poisson	ML	Poisson regression: classical GLM, estimated by maximum likelihood (ML)
		Quasi	“Quasi-Poisson regression”: same mean function, estimated by quasi-ML (QML) or equivalently generalized estimating equations (GEE), inference adjustment via estimated dispersion parameter
		Adjusted	“Adjusted Poisson regression”: same mean function, estimated by QML/GEE, inference adjustment via sandwich covariances
	NB	ML	NB regression: extended GLM, estimated by ML including additional shape parameter
Zero-augmented	Poisson	ML	Zero-inflated Poisson (ZIP), hurdle Poisson
	NB	ML	Zero-inflated NB (ZINB), hurdle NB

Table 1: Overview of various count regression models. The table is usually placed at the top of the page ([t!]), centered (**centering**), has a caption below the table, column headers and captions are in sentence style, and if possible vertical lines should be avoided.

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```
R> data("quine", package = "MASS")
```

and a basic frequency distribution of the response variable is displayed in Figure 1.

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```
R> m_pois <- glm(Days ~ (Eth + Sex + Age + Lrn)^2, data = quine,
+   family = poisson)
```

To account for potential overdispersion we also consider a negative binomial GLM.

```
R> library("MASS")
R> m_nbin <- glm.nb(Days ~ (Eth + Sex + Age + Lrn)^2, data = quine)
```

In a comparison with the BIC the latter model is clearly preferred.

```
R> BIC(m_pois, m_nbin)
```

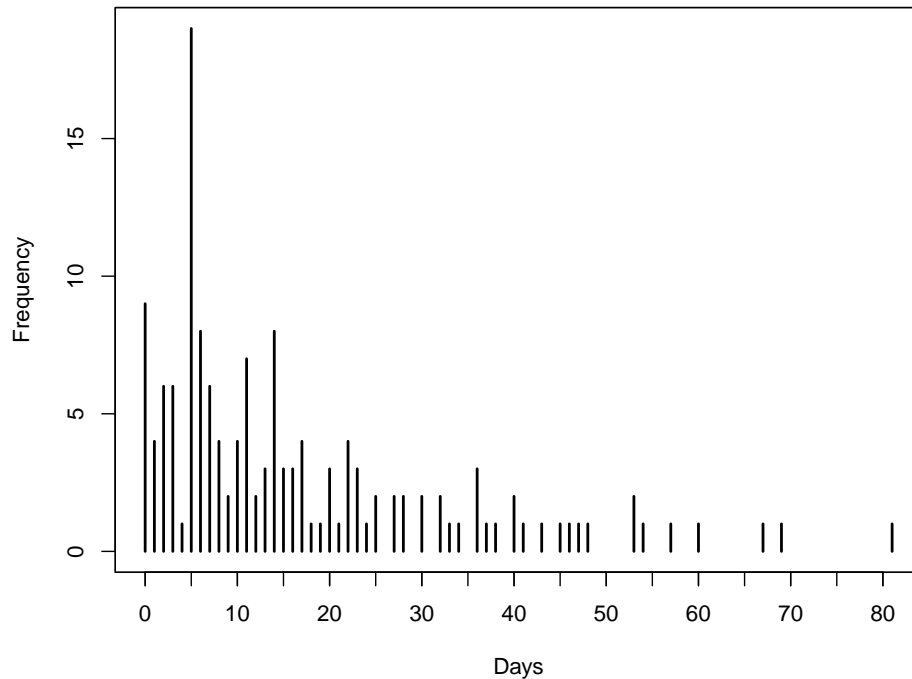


Figure 1: Frequency distribution for number of days absent from school.

```

      df      BIC
m_pois 18 2046.851
m_nbin 19 1157.235

```

Hence, the full summary of that model is shown below.

```
R> summary(m_nbin)
```

Call:

```
glm.nb(formula = Days ~ (Eth + Sex + Age + Lrn)^2, data = quine,
       init.theta = 1.60364105, link = log)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-3.0857	-0.8306	-0.2620	0.4282	2.0898

Coefficients: (1 not defined because of singularities)

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	3.00155	0.33709	8.904	< 2e-16 ***
EthN	-0.24591	0.39135	-0.628	0.52977
SexM	-0.77181	0.38021	-2.030	0.04236 *
AgeF1	-0.02546	0.41615	-0.061	0.95121
AgeF2	-0.54884	0.54393	-1.009	0.31296
AgeF3	-0.25735	0.40558	-0.635	0.52574

LrnSL	0.38919	0.48421	0.804	0.42153
EthN:SexM	0.36240	0.29430	1.231	0.21818
EthN:AgeF1	-0.70000	0.43646	-1.604	0.10876
EthN:AgeF2	-1.23283	0.42962	-2.870	0.00411 **
EthN:AgeF3	0.04721	0.44883	0.105	0.91622
EthN:LrnSL	0.06847	0.34040	0.201	0.84059
SexM:AgeF1	0.02257	0.47360	0.048	0.96198
SexM:AgeF2	1.55330	0.51325	3.026	0.00247 **
SexM:AgeF3	1.25227	0.45539	2.750	0.00596 **
SexM:LrnSL	0.07187	0.40805	0.176	0.86019
AgeF1:LrnSL	-0.43101	0.47948	-0.899	0.36870
AgeF2:LrnSL	0.52074	0.48567	1.072	0.28363
AgeF3:LrnSL	NA	NA	NA	NA

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(Dispersion parameter for Negative Binomial(1.6036) family taken to be 1)

Null deviance: 235.23 on 145 degrees of freedom  
 Residual deviance: 167.53 on 128 degrees of freedom  
 AIC: 1100.5

Number of Fisher Scoring iterations: 1

Theta: 1.604  
 Std. Err.: 0.214

2 x log-likelihood: -1062.546

## 4. Summary and discussion

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## Computational details

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## Acknowledgments

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## References

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