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# RprobitB: Bayes Estimation of Probit Models in R

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#### Abstract

**RprobitB** is an R package for Bayes estimation of multinomial probit models on discrete choice panel data. Its strength lies in modeling heterogenous choices, because it is capable to approximate any underlying mixing distributions through a mixture of normal distributions. The number of latent classes do not need to be prespecified but can be updated within the algorithm on a weight-based strategy. Furthermore, the package provides great numerical stability since it does not require initialization nor approximation or maximization of any likelihood function. In this paper, we show how to estimate (mixed) probit models using **RprobitB** and make a comparision to other R packages that estimate probit models.

Keywords: Mixed multinomial probit models, Bayesian analysis, unobserved heterogeneity.

## 1. Introduction

The introduction is in principle "as usual". However, it should usually embed both the implemented *methods* and the *software* into the respective relevant literature. For the latter both competing and complementary software should be discussed (within the same software environment and beyond), bringing out relative (dis)advantages. All software mentioned should be properly \cite{}d. (See also Appendix B for more details on BibTeX.)

For writing about software JSS requires authors to use the markup \proglang{} (programming languages and large programmable systems), \pkg{} (software packages), \code{} (functions, commands, arguments, etc.). If there is such markup in (sub)section titles (as above), a plain text version has to be provided in the LATEX command as well. Below we also illustrate how abbrevations should be introduced and citation commands can be employed. See the LATEX code for more details.

Modeling count variables is a common task in economics and the social sciences. The classical Poisson regression model for count data is often of limited use in these disciplines because

empirical count data sets typically exhibit overdispersion and/or an excess number of zeros. The former issue can be addressed by extending the plain Poisson regression model in various directions: e.g., using sandwich covariances or estimating an additional dispersion parameter (in a so-called quasi-Poisson model). Another more formal way is to use a negative binomial (NB) regression. All of these models belong to the family of generalized linear models (GLMs). However, although these models typically can capture overdispersion rather well, they are in many applications not sufficient for modeling excess zeros. Since Mullahy (1986) there is increased interest in zero-augmented models that address this issue by a second model component capturing zero counts. An overview of count data models in econometrics, including hurdle and zero-inflated models, is provided in Cameron and Trivedi (2013).

In R (R Core Team 2017), GLMs are provided by the model fitting functions glm() in the stats package and glm.nb() in the MASS package (Venables and Ripley 2002, Chapter 7.4) along with associated methods for diagnostics and inference. The manuscript that this document is based on (Zeileis, Kleiber, and Jackman 2008) then introduced hurdle and zero-inflated count models in the functions hurdle() and zeroinfl() in the pscl package (Jackman 2015). Of course, much more software could be discussed here, including (but not limited to) generalized additive models for count data as available in the R packages mgcv Wood (2006), gamlss (Stasinopoulos and Rigby 2007), or VGAM (Yee 2010).

## 2. Models and software

The basic Poisson regression model for count data is a special case of the GLM framework McCullagh and Nelder (1989). It describes the dependence of a count response variable  $y_i$  (i = 1, ..., n) by assuming a Poisson distribution  $y_i \sim \text{Pois}(\mu_i)$ . The dependence of the conditional mean  $\mathsf{E}[y_i \mid x_i] = \mu_i$  on the regressors  $x_i$  is then specified via a log link and a linear predictor

$$\log(\mu_i) = x_i^{\top} \beta, \tag{1}$$

where the regression coefficients  $\beta$  are estimated by maximum likelihood (ML) using the iterative weighted least squares (IWLS) algorithm.

Note that around the {equation} above there should be no spaces (avoided in the LATEX code by % lines) so that "normal" spacing is used and not a new paragraph started.

R provides a very flexible implementation of the general GLM framework in the function glm() (Chambers and Hastie 1992) in the stats package. Its most important arguments are

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

where formula plus data is the now standard way of specifying regression relationships in R/S introduced in Chambers and Hastie (1992). The remaining arguments in the first line (subset, na.action, weights, and offset) are also standard for setting up formula-based regression models in R/S. The arguments in the second line control aspects specific to GLMs while the arguments in the last line specify which components are returned in the fitted model object (of class 'glm' which inherits from 'lm'). For further arguments to glm() (including

Type	Distribution	Method	Description
GLM	Poisson	ML	Poisson regression: classical GLM, esti-
			mated by maximum likelihood (ML)
		Quasi	"Quasi-Poisson regression": same mean
			function, estimated by quasi-ML (QML)
			or equivalently generalized estimating equa-
			tions (GEE), inference adjustment via esti-
			mated dispersion parameter
		Adjusted	"Adjusted Poisson regression": same mean
			function, estimated by QML/GEE, inference
			adjustment via sandwich covariances
	NB	$\mathrm{ML}$	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.

alternative specifications of starting values) see **?glm**. For estimating a Poisson model **family** = **poisson** has to be specified.

As the synopsis above is a code listing that is not meant to be executed, one can use either the dedicated {Code} environment or a simple {verbatim} environment for this. Again, spaces before and after should be avoided.

Finally, there might be a reference to a {table} such as Table 1. Usually, these are placed at the top of the page ([t!]), centered (\centering), with a caption below the table, column headers and captions in sentence style, and if possible avoiding vertical lines.

#### 3. Illustrations

For a simple illustration of basic Poisson and NB count regression the quine data from the MASS package is used. This provides the number of Days that children were absent from school in Australia in a particular year, along with several covariates that can be employed as regressors. The data can be loaded by

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

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

For code input and output, the style files provide dedicated environments. Either the "agnostic" {CodeInput} and {CodeOutput} can be used or, equivalently, the environments {Sinput} and {Soutput} as produced by Sweave() or knitr when using the render\_sweave() hook. Please make sure that all code is properly spaced, e.g., using y = a + b \* x and not y=a+b\*x. Moreover, code input should use "the usual" command

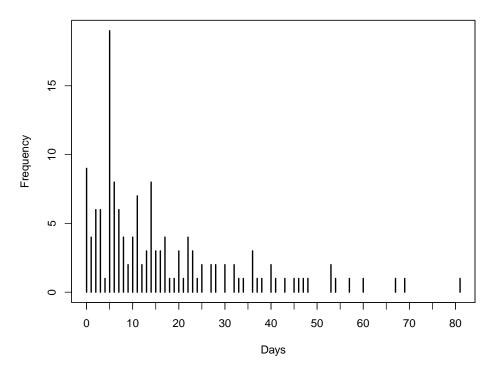


Figure 1: Frequency distribution for number of days absent from school.

prompt in the respective software system. For R code, the prompt "R> " should be used with "+ " as the continuation prompt. Generally, comments within the code chunks should be avoided – and made in the regular LATEX text instead. Finally, empty lines before and after code input/output should be avoided (see above).

As a first model for the quine data, we fit the basic Poisson regression model. (Note that JSS prefers when the second line of code is indented by two spaces.)

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

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)</pre>
```

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

```
R> BIC(m_pois, m_nbin)

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
                    0.38021 -2.030 0.04236 *
          -0.77181
AgeF1
          -0.02546
                    0.41615
                            -0.061 0.95121
AgeF2
          -0.54884
                  0.54393 -1.009 0.31296
          -0.25735
                    0.40558 -0.635 0.52574
AgeF3
LrnSL
          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
                            0.201 0.84059
                    0.34040
SexM:AgeF1
           0.02257
                    0.47360 0.048 0.96198
                  0.51325
                             3.026 0.00247 **
SexM:AgeF2 1.55330
SexM:AgeF3
           1.25227
                  0.45539
                            2.750 0.00596 **
SexM:LrnSL
           0.07187
                    0.40805
                             0.176 0.86019
```

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

NA

0.48567

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

0.47948 -0.899 0.36870

1.072 0.28363

NΑ

NΑ

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

AIC: 1100.5

AgeF3:LrnSL

AgeF1:LrnSL -0.43101

AgeF2:LrnSL 0.52074

Number of Fisher Scoring iterations: 1

NA

Theta: 1.604 Std. Err.: 0.214

2 x log-likelihood: -1062.546

## 4. Summary and discussion

As usual ...

## Computational details

If necessary or useful, information about certain computational details such as version numbers, operating systems, or compilers could be included in an unnumbered section. Also, auxiliary packages (say, for visualizations, maps, tables, ...) that are not cited in the main text can be credited here.

The results in this paper were obtained using R 4.1.0 with the MASS 7.3.54 package. R itself and all packages used are available from the Comprehensive R Archive Network (CRAN) at https://CRAN.R-project.org/.

## Acknowledgments

All acknowledgments (note the AE spelling) should be collected in this unnumbered section before the references. It may contain the usual information about funding and feedback from colleagues/reviewers/etc. Furthermore, information such as relative contributions of the authors may be added here (if any).

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- Software should be properly cited as well. For R packages citation("pkgname") typically provides a good starting point.

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