

Package ‘ZIPBayes’

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Type Package

Title Bayesian Methods in the Analysis of Zero-Inflated Poisson Model

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Description Implementation of zero-inflated Poisson models under Bayesian framework using data augmentation as discussed in Chapter 5 of Zhang (2020) <<https://hdl.handle.net/10012/16378>>. This package is constructed in accommodating four different scenarios: the general scenario, the scenario with measurement error in responses, the external validation scenario, and the internal validation scenario.

License GPL (>= 2)

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ZIPBayes-package

Bayesian Methods in Zero-inflated Poisson Model

Description

Implementation of zero-inflated Poisson (ZIP) model in Bayesian methods with data augmentation strategy. The R package is general to different scenarios, including an ordinary scenario for zero-inflated Poisson Model, the scenario with measurement error in response, the scenario with internal or external validation data available.

Details

This package implemented the zero-inflated Poisson model based on a Bayesian framework. The method is implemented with a Monte Carlo Markov Chain (MCMC) approach with a data augmentation strategy. The package is integrated with C++ to improve the computing speed. It mainly contains four main functions. The function ZIPBayes corresponds to ordinary zero-inflated count data and no measurement error is considered. The function ZIPBayes_MErr considers the case where the response is subject to measurement error as a model by Qihuang Zhang (2020). The function ZIPBayes_Int and ZIPBayes_Ext are corresponding to the case where the internal or external validation data are available, respectively. Other helper functions are also contained in this packages, such as summarizing the trace from the MCMC algorithm, plotting the trace plot, etc.

Author(s)

Qihuang Zhang and Grace Y. Yi.

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datasim

Toy example data - main study only

Description

This data set gives an example data for the illustration of usage of [ZIP](#) and [ZIPMerr](#) function. The dataset contains naivedata and the design matrices for the response model, measurement error model.

Usage

```
data(datasim)
```

Format

A data.frame of 6 columns. "Ystar" refers to the error-prone response. "Y" refers to the true count response. "X1" and "X2" are covariates in the response model. "Xplus" and "Xminus" are the covariates for the measurement error model.

datasimExt*Toy example data - main study and external validation study*

Description

This data set gives an example data for the illustration of usage of [ZIPExt](#) function. The dataset contains a list of main data and external validation data.

Usage

```
data(datasimExt)
```

Format

A list of two data.frames. The first data.frame, named “main”, corresponds to the main data with 6 columns. Same as the [datasim](#), “Ystar” refers to the error-prone response. “Y” refers to the true count response. “X1” and “X2” are covariates in the response model. “Zplus” and “Zplus” are the covariates for the measurement error model. The second data.frame corresponds to the validation data with 5 columns.

datasimInt*Toy example data - main study and internal validation study*

Description

This data set gives an example data for the illustration of usage of [ZIPInt](#) function. The dataset contains a list of main data and internal validation data.

Usage

```
data(datasimInt)
```

Format

A list of two data.frames. The first data.frame, named “main”, corresponds to the main data with 6 columns. Same as the [datasim](#), “Ystar” refers to the error-prone response. “Y” refers to the true count response. “X1” and “X2” are covariates in the response model. “Zplus” and “Zplus” are the covariates for the measurement error model. The second data.frame corresponds to the validation data with 7 columns.

summary.ZIPBayes *Summarizing the trace output from the MCMC algorithm*

Description

This function is a method for **ZIPBayes** object. It summarize the trace output from the main functions into interesting summary statistics, such as mean, median, confidence interval, and highest density region (HDR),

Usage

```
## S3 method for class 'ZIPBayes'

## S3 method for class 'ZIPBayes'
summary(object, burnin = 1, thinperiod = 1, confidence.level = 0.95, ...)
```

Arguments

object	the “ZIPBayes” object gotten from the main function.
burnin	the number of records to be discarded as the early period of MCMC algorithm. Default is 1, meaning the first data point will be discarded when calculating the summary statistics.
thinperiod	the number of period in the thinning period. The results will be picked every this number. Default is 1, meaning no thinning will be done. See details.
confidence.level	the confidence level for the constructed confidence interval. Default is 0.95.
...	other arguments passed to the function.

Details

This function summarizes the tracing results produced by **ZIP**, **ZIPMErr**, **ZIPExt**, and **ZIPInt**. To diminish the influence of the starting values, we generally discard the first portion of each sequence and focus attention on the remaining. The argument **burnin** is set to control the number of steps to be discarded.

Another issue that sometimes arises, once approximate convergence has been reached, is whether to thin the sequences by keeping every k -th simulation draw from each sequence and discarding the rest. The argument **thinperiod** is used to set k here.

Value

ZIPBayes	a list of summary for each data set. "HDR_LB" and "HDR_UB" respectively represents the lower and upper bound of the high density region.
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Author(s)

Qihuang Zhang and Grace Y. Yi.

See Also

[ZIP](#), [ZIPMerr](#), [ZIPExt](#), [ZIPInt](#)

Examples

```
## Please see the example in ZIP() function
```

ZIP

zero-inflated Poisson model

Description

The function implements the MCMC algorithm with data augmentation to estimate the parameters in the zero-inflated Poisson model. The function returns the trace of the sampled parameters in each interaction. To obtain the summary estimation, use `summary()`.

Usage

```
ZIP(Y, Covarmainphi, Covarmainmu,
     betaphi, betamu,
     priorgamma,
     propsigmaphi, propsigmamu = propsigmaphi,
     seed = 1, nmcmc = 500)
```

Arguments

Y	a count vector of length n specifying response in the zero-inflated Poisson model.
Covarmainphi	a $n \times p_1$ dimensional data.frame or matrix of data with respect to the probability component of the zero-inflated Poisson model.
Covarmainmu	a $n \times p_2$ dimensional data.frame or matrix of data with respect to the mean component of the zero-inflated Poisson model.
betaphi	a vector of length p_1 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
betamu	a vector of length p_2 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
priorgamma	a vector of length 2 specifying the two parameters of gamma prior
propsigmaphi	a vector of length p_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the probability component.
propsigmamu	a vector of length p_2 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the mean component.
seed	a numeric value specifying the seed for random generator
nmcmc	a integer specify the number of the generation of MCMC algorithm

Details

The zero-inflated Poisson model involves two components, the probability components and the mean components (Zhang, 2020). Argument Covarmainphi, betaphi, propsigmaphi correspond to the probability component; Covarmainmu, betamu, propsigmamu correspond to the mean component.

Value

BayesResults the list of trace of generated parameters for each component of the models. Data.frame "betaphi_trace" corresponds to the probability component of ZIP response model; "betamu_trace" refers to the mean component of the ZIP response model.

Author(s)

Qihuang Zhang and Grace Y. Yi

References

Zhang, Qihuang. "Inference Methods for Noisy Correlated Responses with Measurement Error." (2020).

See Also

[glm](#)

Examples

```
data(datasim)
set.seed(0)
example_ZIP <- ZIP( Y = datasim$Ystar,
                      Covarmainphi = datasim[,c("intercept","X1")],
                      Covarmainmu = datasim[,c("intercept","X2")],
                      betaphi = c(-0.7,0.7), betamu = c(1,-0.5),
                      priorgamma = rep(1,1), propsigmaphi = c(0.05,0.05),
                      nmcmc = 100)

summary(example_ZIP)
```

ZIPExt

Zero-inflated Poisson model under measurement error and external validation data are available

Description

The function implements the MCMC algorithm with data augmentation to estimate the parameters in the zero-inflated Poisson model while correcting for the measurement error arising from the responses. The function returns the trace of the sampled parameters in each interaction. To obtain the summary estimation, use `summary()`.

Usage

```
ZIPExt (Ystar, Covarmainphi, Covarmainmu, Covarplus, Covarminus,
        Ystarval, Yval, Covarvalplus, Covarvalminus,
        betaphi, betamu, alphaplus, alphaminus,
        Uibound = c(7,11),
        priorgamma, priormu, priorSigma,
        propsigmaphi, propsigmamu = propsigmaphi,
        propsigmaplus = propsigmaphi, propsigmaminus = propsigmaphi,
        seed = 1, nmcmc = 500)
```

Arguments

<code>Ystar</code>	a count vector of length n specifying the error-prone response in the zero-inflated Poisson model.
<code>Covarmainphi</code>	a $n \times p_1$ dimensional data.frame or matrix of the covariate data with respect to the probability component of the zero-inflated Poisson model
<code>Covarmainmu</code>	a $n \times p_2$ dimensional data.frame or matrix of the covariate data with respect to the mean component of the zero-inflated Poisson model
<code>Covarplus</code>	a $n \times q_1$ dimensional data.frame or matrix of the covariate data for the measurement error model of the add-in error process
<code>Covarminus</code>	a $n \times q_2$ dimensional data.frame or matrix of the covariate data for the measurement error model of the leave-out error process
<code>Ystarval</code>	a count vector of length m specifying the error-prone response in the validation data.
<code>Yval</code>	a count vector of length m specifying the precisely measured response in the validation data.
<code>Covarvalplus</code>	a $m \times q_1$ dimensional data.frame or matrix of the covariate for validation data of the add-in error process
<code>Covarvalminus</code>	a $m \times q_2$ dimensional data.frame or matrix of the covariate for validation data of the leave-out error process
<code>betaphi</code>	a vector of length p_1 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
<code>betamu</code>	a vector of length p_2 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
<code>alphaplus</code>	a vector of length q_1 specifying the initial values of the parameters for the measurement error model of the add-in error process
<code>alphaminus</code>	a vector of length q_2 specifying the initial values of the parameters in the probability component of the leave-out error process
<code>Uibound</code>	a vector of length 2 specifying the maximum number of the count in the inverse sampling method
<code>priorgamma</code>	a vector of length 2 specifying the two parameters of gamma prior
<code>priormu</code>	a vector of length q_2 specifying the mean vector of the normal prior for the measurement error model of the leave-out error process

priorSigma	a vector of length q_2 specifying the standard errors of the normal prior for the measurement error model of the leave-out error process
propsigmaphi	a vector of length p_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the probability component.
propsigmamu	a vector of length p_2 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the mean component.
propsigmaplus	a vector of length q_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the add-in error process.
propsigmaminus	a vector of length q_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the leave-out error process.
seed	a numeric value specifying the seed for random generator
nmcmc	a integer specify the number of the generation of MCMC algorithm

Details

Comparing to the ZIPMerr function, this function has an addition component – validation data. Here, the argument “Ystarval”, “Yval”, “Covarvalplus”, “Covarvalminus”, are new for the sceanrio with external validation.

Value

BayesResults	the list of trace of generated parameters for each component of the models. Data frame “betaphi_trace” corresponds to the probability component of ZIP response model; “betamu_trace” refers to the mean component of the ZIP response model. Data frames “alphaplus_trace” and “alphaminus_trace”, respectively, correspond to the add-in error and leave-out error process in the measurement error model.
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Author(s)

Qihuang Zhang and Grace Y. Yi

References

Zhang, Qihuang. “Inference Methods for Noisy Correlated Responses with Measurement Error.” (2020).

See Also

[glm](#)

Examples

```
## load data
data(datasimExt)
set.seed(0)
example_ZIP_Ext <- ZIPExt (Ystar = datasimExt$main$Ystar,
                           Covarmainphi = datasimExt$main[,c("intercept","X1")],
```

```

Covarmainmu = datasimExt$main[,c("intercept","X2")],
Covarplus = datasimExt$main[,c("intercept","Zplus")],
Covarminus = datasimExt$main[,c("intercept","Zminus")],
Ystarval = datasimExt$validation$Ystar,
Yval = datasimExt$validation$Y,
Covarvalplus = datasimExt$validation[,3:4],
Covarvalminus = datasimExt$validation[,3:4],
betaphi = c(0.7,-0.7), betamu = c(1,-1.5),
alphaplus = c(0,0), alphaminus=c(0,0),
priorgamma = c(0.001,0.001), priormu = c(0,0),
priorSigma = c(1,1), propsigmaphi = c(0.05,0.05),
nmcmc = 10)

summary(example_ZIP_Ext)

```

ZIPInt

Zero-inflated Poisson model under measurement error and internal validation data are available

Description

The function implements the MCMC algorithm with data augmentation to estimate the parameters in the zero-inflated Poisson model while correcting for the measurement error arising from the responses. The function returns the trace of the sampled parameters in each interaction. To obtain the summary estimation, use `summary()`.

Usage

```

ZIPInt (Ystar, Covarmainphi, Covarmainmu, Covarplus, Covarminus,
        Ystarval, Yval, Covarvalmainphi, Covarvalmainmu, Covarvalplus, Covarvalminus,
        betaphi, betamu, alphaplus, alphaminus,
        Uibound = c(7,11),
        priorgamma, priormu, priorSigma,
        propsigmaphi, propsigmamu = propsigmaphi,
        propsigmaphi, propsigmaphi, propsigmaminus = propsigmaphi,
        seed = 1, nmcmc = 500)

```

Arguments

- | | |
|---------------------------|--|
| <code>Ystar</code> | a count vector of length n specifying the error-prone response in the zero-inflated Poisson model. |
| <code>Covarmainphi</code> | a $n \times p_1$ dimensional data.frame or matrix of the covariate data with respect to the probability component of the zero-inflated Poisson model |
| <code>Covarmainmu</code> | a $n \times p_2$ dimensional data.frame or matrix of the covariate data with respect to the mean component of the zero-inflated Poisson model |
| <code>Covarplus</code> | a $n \times q_1$ dimensional data.frame or matrix of the covariate data for the measurement error model of the add-in error process |

Covarminus	a $n \times q_2$ dimensional data.frame or matrix of the covariate data for the measurement error model of the leave-out error process
Ystarval	a count vector of length m specifying the error-prone response in the validation data.
Yval	a count vector of length m specifying the precisely measured response in the validation data.
Covarvalmainphi	a $m \times p_1$ dimensional data.frame or matrix of the covariate for validation data with respect to the probability component of ZIP model
Covarvalmainmu	a $m \times p_2$ dimensional data.frame or matrix of the covariate for validation data with respect to the mean component of ZIP model
Covarvalplus	a $m \times q_1$ dimensional data.frame or matrix of the covariate for validation data of the add-in error process
Covarvalminus	a $m \times q_2$ dimensional data.frame or matrix of the covariate for validation data of the leave-out error process
betaphi	a vector of length p_1 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
betamu	a vector of length p_2 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
alphaplus	a vector of length q_1 specifying the initial values of the parameters for the measurement error model of the add-in error process
alphaminus	a vector of length q_2 specifying the initial values of the parameters in the probability component of the leave-out error process
Uibound	a vector of length 2 specifying the maximum number of the count in the inverse sampling method
priorgamma	a vector of length 2 specifying the two parameters of gamma prior
priormu	a vector of length q_2 specifying the mean vector of the normal prior for the measurement error model of the leave-out error process
priorSigma	a vector of length q_2 specifying the standard errors of the normal prior for the measurement error model of the leave-out error process
propsigmaphi	a vector of length p_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the probability component.
propsigmamu	a vector of length p_2 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the mean component.
propsigmaplus	a vector of length q_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the add-in error process.
propsigmaminus	a vector of length q_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the leave-out error process.
seed	a numeric value specifying the seed for random generator
nmcmc	a integer specify the number of the generation of MCMC algorithm

Details

Comparing to the ZIPExt function for the external validation study, this function has an addition component – covariates of response model in the validation data. Here, the argument “Covarvalmainphi” and “Covarvalmainmu” are new for the scenario with external validation.

Value

`BayesResults` the list of trace of generated parameters for each component of the models. Data frame "betaphi_trace" corresponds to the probability component of ZIP response model; "betamu_trace" refers to the mean component of the ZIP response model. Data frames "alphaplus_trace" and "alphaminus_trace", respectively, correspond to the add-in error and leave-out error process in the measurement error model.

Author(s)

Qihuang Zhang and Grace Y. Yi

References

Zhang, Qihuang. "Inference Methods for Noisy Correlated Responses with Measurement Error." (2020).

See Also

[glm](#)

Examples

```
data(datasimInt)
set.seed(0)
result <- ZIPInt(Ystar = datasimInt$main$Ystar,
                  Covarvalmainphi = datasimInt$val[,c("intercept", "X1")],
                  Covarvalmainmu = datasimInt$val[,c("intercept", "X2")],
                  Covarvalplus = datasimInt$val[,c("intercept", "Zplus")],
                  Covarvalminus = datasimInt$val[,c("intercept", "Zminus")],
                  Ystarval = datasimInt$main$Ystar,
                  Yval = datasimInt$main$Y,
                  Covarvalmainphi = datasimInt$val[,c("intercept", "X1")],
                  Covarvalmainmu = datasimInt$val[,c("intercept", "X2")],
                  Covarvalplus = datasimInt$val[,c("intercept", "Zplus")],
                  Covarvalminus = datasimInt$val[,c("intercept", "Zminus")],
                  betamu = c(-0.5,0.5), betaphi = c(0.5,0),
                  alphaplus = c(0,0), alphaminus= c(0,0),
                  priorgamma = c(1,1), priormu = c(0,0), priorSigma = c(1,1),
                  propsigmaphi = c(0.05,0.05), nmcmc = 10)

summary(result)
```

ZIPMerr*Zero-inflated Poisson model with measurement error*

Description

The function implements the MCMC algorithm with data augmentation to estimate the parameters in the zero-inflated Poisson model while correcting for the measurement error arising from the responses. The function returns the trace of the sampled parameters in each iteration. To obtain the summary estimation, use `summary()`.

Usage

```
ZIPMerr (Ystar, Covarmainphi, Covarmainmu, Covarplus, Covarminus,
         betaphi, betamu, alphaplus, alphaminus,
         Uibound = c(7,11),
         priorgamma, priormu, priorSigma, propsigmaphi, propsigmamu = propsigmaphi,
         propsigmaplus = propsigmaphi, propsigmaminus = propsigmaphi,
         seed = 1, nmcmc = 500)
```

Arguments

<code>Ystar</code>	a count vector of length n specifying the error-prone response in the zero-inflated Poisson model.
<code>Covarmainphi</code>	a $n \times p_1$ dimensional data.frame or matrix of the covariate data with respect to the probability component of the zero-inflated Poisson model
<code>Covarmainmu</code>	a $n \times p_2$ dimensional data.frame or matrix of the covariate data with respect to the mean component of the zero-inflated Poisson model
<code>Covarplus</code>	a $n \times q_1$ dimensional data.frame or matrix of the covariate data for the measurement error model of the add-in error process
<code>Covarminus</code>	a $n \times q_1$ dimensional data.frame or matrix of the covariate data for the measurement error model of the leave-out error process
<code>betaphi</code>	a vector of length p_1 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
<code>betamu</code>	a vector of length p_2 specifying the initial values of the parameters in the probability component of the zero-inflated Poisson model
<code>alphaplus</code>	a vector of length q_1 specifying the initial values of the parameters for the measurement error model of the add-in error process
<code>alphaminus</code>	a vector of length q_2 specifying the initial values of the parameters in the probability component of the leave-out error process
<code>Uibound</code>	a vector of length 2 specifying the maximum number of the count in the inverse sampling method
<code>priorgamma</code>	a vector of length 2 specifying the two parameters of gamma prior
<code>priormu</code>	a vector of length q_2 specifying the mean vector of the normal prior for the measurement error model of the leave-out error process

priorSigma	a vector of length q_2 specifying the standard errors of the normal prior for the measurement error model of the leave-out error process
propsigmaphi	a vector of length p_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the probability component.
propsigmamu	a vector of length p_2 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the mean component.
propsigmaplus	a vector of length q_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the add-in error process.
propsigmaminus	a vector of length q_1 specifying the standard error of the Gaussian proposal distribution for the parameters corresponds to the leave-out error process.
seed	a numeric value specifying the seed for random generator
nmcmc	a integer specify the number of the generation of MCMC algorithm

Details

The zero-inflated Poisson model involves two components, the probability components and the mean components (Zhang, 2020). Although there are might arguments involved in the functions, they can be summarized to four sources in the model. The response model (zero-inflated Poisson model) involves two components: the probability component and the mean count component. The measurement error models contains two process: the add-in process and leave-out process. The arguments end with "-phi" corresponds to the probability component of the response model. The arguments end with "-mu" corresponds to the mean component of the response model. The arguments end with "-plus" corresponds to the add-in error process in the measurement error model. The arguments end with "-minus" corresponds to the leave-out process of the measurement error model.

Value

BayesResults	the list of trace of generated parameters for each component of the models. Data frame "betaphi_trace" corresponds to the probability component of ZIP response model; "betamu_trace" refers to the mean component of the ZIP response model. Data frames "alphaplus_trace" and "alphaminus_trace", respectively, correspond to the add-in error and leave-out error process in the measurement error model.
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Author(s)

Qihuang Zhang and Grace Y. Yi

References

Zhang, Qihuang. "Inference Methods for Noisy Correlated Responses with Measurement Error." (2020).

See Also

[glm](#)

Examples

```

## load data
data(datasim)
set.seed(0)
example_ZIP_MErr <- ZIPMERR (Ystar = datasim$Ystar,
                               Covarmainphi = datasim[,c("intercept","X1")],
                               Covarmainmu = datasim[,c("intercept","X2")],
                               Covarplus = datasim[,c("intercept","Xplus")],
                               Covarminus = datasim[,c("intercept","Xminus")],
                               betaphi = c(0.7,-0.7), betamu = c(1,-1.5),
                               alphaplus = c(0,0), alphaminus=c(0,0),
                               priorgamma = c(0.001,0.001), priormu = c(0,0),
                               priorSigma = c(1,1), propsigmaphi = c(0.05,0.05),
                               nmcmc = 10)

summary(example_ZIP_MErr)

```

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