

Package ‘svyVGAM’

September 30, 2020

Title Design-Based Inference in Vector Generalised Linear Models

Version 0.222-2

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Description

Provides design-based inference for the wide range of parametric models in the 'VGAM' package.

Imports stats, methods

Depends VGAM, survey, R (>= 3.5.0)

Suggests pscl, knitr, markdown, rmarkdown

VignetteBuilder knitr

Encoding UTF-8

License GPL-3

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NeedsCompilation no

Repository CRAN

Date/Publication 2020-09-30 08:50:02 UTC

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nhanes_sxq	<i>Data from NHANES: number of sex partners</i>
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Description

These data are from the NHANES 2003-2004 survey in the US. They provide an example of overdispersed count data that motivates a two-component zero-inflation model

Usage

```
data("nhanes_sxq")
```

Format

A data frame with 2992 observations on the following 7 variables.

SDMVPSU Primary Sampling Unit

SDMVSTRA stratum

WTINT2YR weights

malepartners lifetime number of male sexual partners

RIDAGEYR age in years

DMD EDUC level of education: 1=less than high school, 2=high school, 3=more than high school, 7=refused

RIDRETH1 Race/ethnicity: 1=Mexican American, 2=Other Hispanic, 4=non-Hispanic White, 5=non-Hispanic Black, 5=Other

Source

NHANES files demo_c.xpt and sxq_c.xpt

See Also

Construction of the data set is described by <https://notstatschat.rbind.io/2015/05/26/zero-inflated-poisson-fr>

Examples

```
data(nhanes_sxq)
nhdes = svydesign(id=~SDMVPSU, strat=~SDMVSTRA, weights=~WTINT2YR, nest=TRUE, data=nhanes_sxq)
svy_vglm(malepartners~RIDAGEYR+factor(RIDRETH1)+DMD EDUC, zipoisson(), design=nhdes, crit = "coef")
```

 svy_vglm

Design-based inference for vector generalised linear models

Description

This function provides design-based (survey) inference for Thomas Yee's vector generalised linear models. It works by calling `vglm` with sampling weights, and then either using resampling (replicate weights) or extracting the influence functions and using a Horvitz-Thompson-type sandwich estimator.

Usage

```
svy_vglm(formula, family, design, ...)
```

Arguments

formula	Model formula, as for vglm
family	Model family, as for vglm
design	Survey design object
...	Other arguments to pass to vglm

Value

An S3 object of class `svy_glm` with `print`, `coef` and `vcov` methods, containing the design in the `design` component and a fitted `vglm` object in the `fit` component.

See Also

[nhanes_sxq](#)
[vglm](#)
[svydesign](#) [svrepdesign](#)

Examples

```
data(api)
dclus2<-svydesign(id=~dnum+snum, fpc=~fpc1+fpc2, data=apiclus2)

## Ordinary Gaussian regression
m1<-svyglm(api00~api99+mobility+ell, design=dclus2,family=gaussian)
## same model, but with the variance as a second parameter
m2<-svy_vglm(api00~api99+mobility+ell, design=dclus2,family=uninormal())
m1
m2
SE(m1)
SE(m2)

summary(m1)
summary(m2)

## Proportional odds model
dclus2<-update(dclus2, mealcat=as.ordered(cut(meals,c(0,25,50,75,100))))
a<-svyolr(mealcat~avg.ed+mobility+stype, design=dclus2)
b<-svy_vglm(mealcat~avg.ed+mobility+stype, design=dclus2, family=propodds())
a
b
SE(a)
SE(b) #not identical, because svyolr() uses approximate Hessian

## Zero-inflated Poisson
data(nhanes_sxq)
nhdes = svydesign(id=~SDMVPSU, strat=~SDMVSTRA, weights=~WTINT2YR,
  nest=TRUE, data=nhanes_sxq)

sv1<-svy_vglm(malepartners~RIDAGEYR+factor(RIDRETH1)+DMDEDUC,
```

```
      zipoisson(), design=nhdes, crit = "coef")
sv1
summary(sv1)

## Multinomial
## Reference group (non-Hispanic White) average older and more educated
## so coefficients are negative
mult_eth<- svy_vglm(RIDRETH1~RIDAGEYR+DMDEDUC,
  family=multinomial(refLevel=3), design=nhdes)

## separate logistic regressions are close but not identical
two_eth<-svyglm(I(RIDRETH1==1)~RIDAGEYR+DMDEDUC, family=quasibinomial,
  design=subset(nhdes, RIDRETH1 %in% c(1,3)))

summary(mult_eth)
summary(two_eth)
```

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