

Package: svyVGAM (via r-universe)

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Title Design-Based Inference in Vector Generalised Linear Models

Version 1.2-1

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Description Provides inference based on the survey package 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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Repository <https://tstumley.r-universe.dev>

RemoteUrl <https://github.com/tstumley/svyvgam>

RemoteRef HEAD

RemoteSha 7eed207169016d00cbe52a54ad514c73b30b6495

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`nhanes_sxq`*Data from NHANES: number of sex partners*

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

DMDEDUC 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)+DMDEDUC, zipoisson(), design=nhdes, crit = "coef")
```

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

<code>formula</code>	Model formula, as for <code>vglm</code>
<code>family</code>	Model family, as for <code>vglm</code>
<code>design</code>	Survey design object
<code>...</code>	Other arguments to pass to <code>vglm</code>

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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