Robust inference for generalized linear models by flipscores approach

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A linear regression model is defined as

$$Y = X\beta + Z\gamma + \varepsilon$$

where

- Y is the outcome
- X, Z are observed covariates
- β,γ are regression coefficients, β is of direct interest
- $\varepsilon \sim N(0, \sigma^2 I)$ is an error term
- \bullet the variance σ^2 is assumed common among all the units
- Consider dim $(\beta) = 1$ and dim $(\gamma) \ge 1$

We want to test $H_0: \beta = 0$ against a one or two-sided alternative

- β is the parameter of interest
- γ,σ are nuisance parameters, not of direct interest but we have to estimate them

What happens if we ignore some existing heteroscedasticity?



Some simulated data with

- $y_i \sim N(\mu_i, \sigma_i^2)$
- $\mu_i = x_i\beta + z_i\gamma$
- $cor(x_i, z_i) = 0.5$
- β = 0
- $\gamma = 1$
- $\sigma_i^2 = 4x_i^2$.

We fit a linear model assuming common variance, testing H_0 : $\beta = 0$ with significance level $\alpha = 0.05$



Sample size	Proportion of rejection
25	0.20
50	0.21
100	0.21
200	0.21
500	0.22
1000	0.21

Much higher than 0.05, we reject too often (no type I error control!)



Generalized linear models are a flexible tool introduced to extend linear regression models. Some examples are

- normal regression with logarithmic link
- poisson regression
- Iogistic regression

Usually we have a strong assumption on the variance structure (e.g. homoscedastic Normal model, Poisson model, \dots).

Further problem: when the model variance is not constant it is difficult to check the validity of the assumptions

The regression model is

$$g(\mu_i) = \eta_i = x_i\beta + z_i\gamma.$$

Aim: we consider univariate test of the form

$$H_0: \beta = 0$$

against a one or two-sided alternative.

We want to build a test robust against variance misspecification



 $\ensuremath{\textbf{Sign-flip}}$ tests offer an alternative way to do hypothesis testing. Usually they

- require less assumptions (semi-parametric tests)
- converge to the parametric counterpart (when it exists)
- have exact control of type I error



What are sign-flips?

Suppose we have a sample of *n* observations. Sign-flips are *n*-dimensional vectors of 1 and -1. Example, n = 6:

$$I = F_1 = (1, 1, 1, 1, 1, 1)$$
$$F_2 = (1, 1, 1, 1, 1, -1)$$
$$\vdots$$
$$F_{64} = (-1, -1, -1, -1, -1, -1)$$

In general the total amount is 2^n different flips.

The idea is to use a conditional (flipping) distribution of the data. What does it mean?

- Let T(I) be any observed test statistic.
- Call *T*(*F*) a flipped test statistic. It is obtained by multiplying the data (or other appropriate quantities) by a sign-flip *F*.
- We have a flipping distribution with a total of 2ⁿ test statistics.
- We can perform valid hypothesis testing if

$$T(I) \stackrel{d}{=} T(F)$$
 (equality in distribution)

for all sign-flips, when H_0 is true

- We observe a sample of independent observations y_1, \ldots, y_n , with $y_i \sim N(\mu, \sigma_i)$
- We test $H_0: \mu = 0$ vs $H_1: \mu > 0$, significance level of α
- When H_0 is true, $y_i \stackrel{d}{=} -y_i$ (equality in distribution).
- Use the test statistic $T(I) = \sum_{i=1}^{n} y_i$
- The flipped test statistic is $T(F) = \sum_{i=1}^{n} f_i y_i$
- We have $T(I) \stackrel{d}{=} T(F)$



- Call $G = 2^n$ the total amount of transformations.
- We order them

$$T^{(1)}(F) \leq \cdots \leq T^{(G)}(F)$$

- We reject H₀ if T(I) > T^{(⌈(1-α)·G⌉)}(F) (which is the 1 − α quantile of the flipping distribution)
- Exact control of type I error



We observe a sample of 6 observations $(2^6 = 64)$

$$Y = (-1.63, 1.61, 0.13, 0.66, 0.01, -0.65)$$

$$T(I) = -1.63 + 1.61 + 0.13 + 0.66 + 0.01 - 0.65 = 0.13$$

$$T(F_2) = 1.63 + 1.61 + 0.13 + 0.66 + 0.01 - 0.65 = 3.39$$

$$T(F_3) = 1.63 - 1.61 + 0.13 + 0.66 + 0.01 - 0.65 = 0.17$$

 $T(F_{64}) = 1.63 - 1.61 - 0.13 - 0.66 - 0.01 + 0.65 = -0.13$

We reject H_0 if $T(I) > T^{(61)}(F)$

. . .

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How to apply the idea of sign-flip tests for GLMs?

• The outcome Y cannot be used. Under $H_0: \beta = 0$

$$\mu_i = g^{-1} \left(0 + z_i \gamma \right)$$

In general

$$\mu_i \neq -\mu_i$$

$$\implies y_i \neq -y_i$$

$$\implies T(I) \neq T(F$$



New proposal: use the effective score. It is defined as

$$T(F) = n^{-1/2} F(S_{\beta} - \mathcal{I}_{\beta\gamma} \mathcal{I}_{\gamma\gamma}^{-1} S_{\gamma}) = n^{-1/2} \sum_{i=1}^{n} f_{i} \nu_{i}^{*} \Big|_{\beta = 0, \gamma = \hat{\gamma}}$$

We have

 $\mathbb{E}[T(I)] = \mathbb{E}[T(F)] = 0, \quad \mathbb{V}[T(F)] \xrightarrow{n \to \infty} \mathbb{V}[T(I)]$



Easy solution to improve the convergence of the test statistic. Standardized test statistic

$$T_{s}(F) = T(F)/\mathbb{V}(T(F))^{1/2}.$$

We have

 $\mathbb{E}[T_s(I)] = \mathbb{E}[T_s(F)] = 0, \quad \mathbb{V}[T_s(I)] = \mathbb{V}[T_s(F)] = 1$

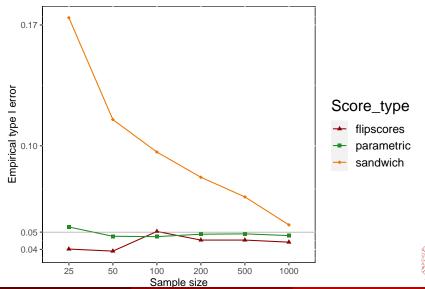


Example we simulate *n* observations from the model $y_i \sim Poisson(\mu_i)$

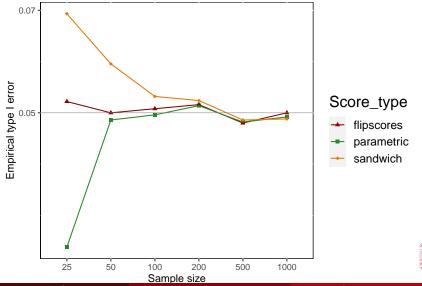
- $\log(\mu_i) = x_i\beta + z_i\gamma$
- $(\beta, \gamma) = (0, 1, 1, 1)$
- $cor(x_i, z_i) = (0.5, 0.1, 0.1)$

We have two competitors: the **standard parametric test** and the **sandwich estimator**. See some simulations!





Simulation: Logit model



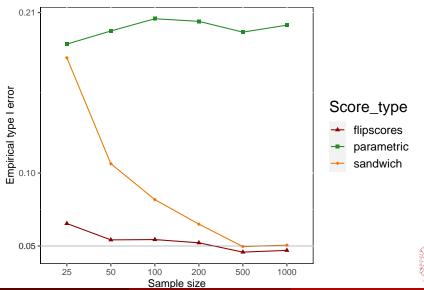
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Linear model with ignored heteroscedasticity. Some simulated data with

- $\beta = 0$
- $\gamma = (1, 1, 1)$
- $\sigma_i^2 = 4x_i^2$
- $cor(x_i, z_i) = (0.5, 0.1, 0.1)$



Ignored Heteroscedastic Normal



Poisson distribution

- $\mathcal{Y} = \mathbb{N}$
- $\mathbb{E}[y_i] = \mu_i$
- $\mathbb{V}[y_i] = \mu_i$

Negative binomial distribution

•
$$\mathcal{Y} = \mathbb{N}$$

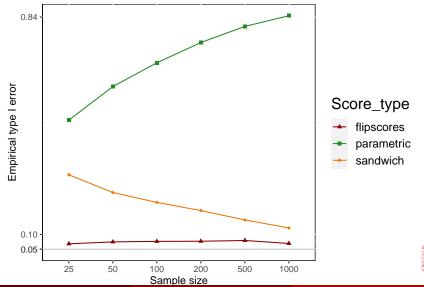
•
$$\mathbb{E}[y_i] = \mu_i$$

•
$$\mathbb{V}[y_i] = \mu_i \left(1 + \phi \mu_i\right)$$

We fit a Poisson regression model while the true distribution is Negative binomial. We set $\phi=1$



True Negative binomial, fitted Poisson



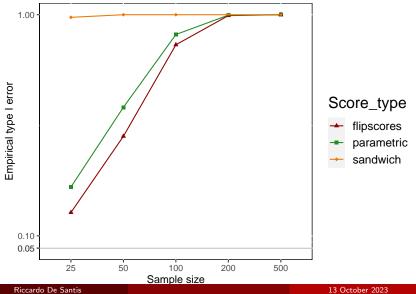
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If the test is able to control the type I error, it should have good power

For a meaningful comparison we fit a correctly specified Poisson model setting $\beta = 0.3$, while we test H_0 : $\beta = 0$



Power comparison



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With n = 1000 and 4 covariates the computational time is ≈ 4.2 seconds

> summary(mod)

```
Call:
flipscores(formula = Y \sim X + Z, family = poisson(link = "log"),
   score_type = "standardized")
Deviance Residuals:
   Min
            10 Median
                             30
                                     Max
-1.0842 -0.9145 -0.7003 0.2292 2.3617
Coefficients:
           Estimate Score Std. Error z value eff size Pr(>|z|)
(Intercept) -1.0072 -2.5093 0.6527 -3.8442 -0.630 0.0012 **
           0.7292 1.3630 0.9683 1.4076 0.389 0.2058
Х
            -1.0395 -1.1693 0.5107 -2.2896 -0.465 0.0372 *
z
signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



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