The paper solves the bias in data arising from non-random testing of entire population for Covid-19. A two-equation model overcomes the bias by using the inverse Mills ratio and improves forecasts of new deaths in all nine out of nine weekly out-of-sample comparisons. We identify the political party of the governors of states with desirable negative and undesirable positive trends.

```
library(sampleSelection)
library(margins)
library(dplyr)
##Pull Data
Cumulative Deaths=read.csv("Cumulative Deaths.csv")
Cumulative Infections=read.csv("Cumulative Infections.csv")
Covid=read.csv("Covid 0615.csv")
##clean data
Covid$State.Abbreviation<-Covid$STA
Final Dataset<-merge(Covid, Cumulative Deaths, by="State. Abbreviation")
Data<-merge (Final Dataset, Cumulative Infections, by="State.Abbreviation")
2020/06/15: First Step Probit Model
myprobit<-glm(TI~HES+CPT+UI+HR+HI, family=binomial(link="probit"),data=Data)</pre>
##TI is Testing Indicator (i.e. 1 for tested and 0 for not tested)
##HES is Hospital Employee Share
##CPT is portion of population that Commutes on Public Transit
##UI in Uninsured Population
##HR is Hypertension Rate (one of the leading comorbidities)
#HI is Household Income
summary(myprobit)
##
## Call:
## glm(formula = TI ~ HES + CPT + UI + HR + HI, family = binomial(link =
"probit"),
##
     data = Data)
##
## Deviance Residuals:
     Min 1Q Median 3Q
                                      Max
## -0.7296 -0.3849 -0.3553 -0.3376 2.5180
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -2.4319390 0.0636110 -38.231 < 2e-16 ***
             ## HES
             0.0193450 0.0004345 44.523 < 2e-16 ***
## CPT
            -0.0070335 0.0009966 -7.058 1.69e-12 ***
## UI
             ## HR
              0.0021614 0.0004608 4.690 2.73e-06 ***
## HI
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 254187 on 499999 degrees of freedom
## Residual deviance: 250590 on 499994 degrees of freedom
## AIC: 250602
##
## Number of Fisher Scoring iterations: 5
2020/06/15: Probit Model Marginal Effects
```

```
#Marginal Effects of probit model
effects pro=margins(myprobit) #print(effects pro)
summary(effects pro)
  CPT 0.0025685341 5.801388e-05
                                      44.274475 0.000000e+00
                                                                  0.0024548290
0.0026822392
  HES 0.0088046672 1.084429e-03 8.119170 4.693813e-16 0.0066792246
0.0109301098
3 HI 0.0002869787 6.119577e-05 4.689519 2.738481e-06 0.0001670372
0.0004069203
4 HR 0.0026358250 1.465264e-04
                                      17.988742 2.387156e-72 0.0023486386
0.0029230114
5 UI -0.0009338730 1.323531e-04 -7.055923 1.714593e-12
-0.0011932802 -0.0006744657
5 rows
2020/06/15: Extract inverse Mills ratio from probit model
##Compute inverse Mills ratio from probit model usine 'invMillsRatio' command
from the 'sampleSelection' package.
IMR=invMillsRatio(myprobit,all=FALSE)
Data$IMR<-invMillsRatio(myprobit)$IMR1</pre>
##create table of IMR values by state
IMR.Index=cbind(as.character(Data$State.Abbreviation),Data$IMR)
IMR table=distinct(data.frame(IMR.Index))
2020/06/15: Second Step Heckman Equation (i.e. with IMR)
##GLM with Poisson distribution to estimate cumulative deaths based on the log
of cumulative infections from the previous week, as well as the IMR.
##Use 'glm' command with 'family=poisson(link="log")'
##Only ran the model on individuals who were tested (i.e. <math>TI = 1).
##CD is "Cumulative Deaths"
##CI is "Cumulative Infections"
glm IMR 0615 CD<-glm(X20200615 CD~log(X20200608 CI)
+IMR, family=poisson(link="log"),
                    data=subset(Data,TI==1))
fig-out-of-sample-all
#Compute fitted values for each state
Tested Individuals<-subset(Data, TI==1)</pre>
Fitted CD IMR=fitted(glm IMR 0615 CD)
Fitted_CD=fitted(glm_0615_CD)
Results Table=cbind(as.character(Tested Individuals$State.Abbreviation),Tested
Individuals$X20200622 CI,
              Tested Individuals$X20200622 CD, Tested Individuals$
X20200615 CD, Fitted CD IMR, Fitted CD,
              Tested Individuals$IMR)
#Pull fitted value for each state
Final Results=distinct(data.frame(Results Table))
View(Final_Results)
Assessing Nation-wide Forecast
```

```
#Compare fitted values with in-sample (2020/06/15) and out-of-sample
(2020/06/22) actual deaths
Actual Infections Out Sample=sum(as.numeric(Final Results[,2]))
Actual Deaths Out Sample=sum(as.numeric(Final Results[,3]))
Predict Deaths IMR=sum(as.numeric(Final Results$Fitted CD IMR))
Predict Deaths no IMR=sum(as.numeric(Final Results$Fitted CD))
Out of sample per diff CD IMR=(Predict Deaths IMR-Actual
Deaths Out Sample) / Actual Deaths Out Sample
Out of sample per diff CD no IMR=(Predict Deaths no IMR-
Actual Deaths Out Sample)/Actual Deaths Out Sample
Actual Deaths In Sample=sum(as.numeric(Final Results[,4]))
In sample per diff CD IMR=(Predict Deaths IMR-Actual Deaths In Sample)/Actual
Deaths In Sample
In sample per diff CD no IMR=(Predict Deaths no IMR-Actual
Deaths In Sample)/Actual Deaths In Sample
print(rbind(Actual Deaths In Sample, In sample per diff CD
IMR, In sample per diff CD no IMR))
##
                                        [,1]
## Actual Deaths In Sample
                               1.098220e+05
## In sample per diff CD IMR
                               4.064085e-03
\#\# In sample per diff CD no IMR -4.705036e-02
print(rbind(Actual Infections Out Sample, Actual Deaths Out
Sample, Predict Deaths no IMR, Out of sample per diff CD IMR,
Out of sample per diff CD no IMR))
                                            [,1]
## Actual_Infections_Out_Sample 2.290489e+06
## Actual_Deaths_Out_Sample
## Predict_Deaths_no_IMR
                                   1.139440e+05
                                   1.046548e+05
## Out_of_sample_per_diff_CD_IMR -3.225860e-02
\#\# Out of sample per diff CD no IMR -8.152394e-02
In all nine cases using IMR bias correction improves out-of-sample forecasts of
Covid-19 deaths in US as a whole
cbind(Week,IMR Perc Diff,Without IMR Perc Diff)
## Week IMR Perc Diff Without IMR Perc Diff
## [1,] 1 -25.27
                                         -26.96
## [2,] 2
                   -21.61
                                        -22.50
## [3,] 3
                   -17.66
                                         -18.73
## [4,] 4
                  -12.14
                                         -13.78
## [5,] 5
                   -9.38
                                        -11.28
## [6,] 6
## [7,] 7
                   -7.25
                                          -9.70
                -5.53
-3.94
                                          -9.01
## [8,] 8
                                          -8.03
```