```
library(pscl)
model1 = zeroinfl(Infections ~ Swimmer +Location +Age +Sex | Swimmer +
                   Location+ Age+ Sex, dist = "negbin", data = earinfection)
summary(model1)
summary(model2 <- glm(Infections ~ Swimmer + Location + Age + Sex,</pre>
                         family = poisson, data = earinfection))
vuong(model2, model1)
Call:
zeroinfl(formula = Infections ~ Swimmer + Location + Age + Sex | Swimmer + Location + Age + Sex, data = earinfection, dist = "negbin")
 Pearson residuals:
 Min 1Q Median 3Q Max
-0.7741 -0.6467 -0.4308 0.3747 6.1088
Count model coefficients (negbin with log link):
Estimate Std. Error z value Pr(>|z|)
(Intercept) 2.10576 0.65143 3.232 0.001227 **
                                      -3.443 0.000576 ***
 Swimmer
              -0.67532
                            0.19616
               -0.06729
                            0.23856
                                      -0.282 0.777895
 Location
                            0.02285
                                       -0.744 0.456951
              -0.01700
 Aae
                                       -0.690 0.490321
               -0.15246
Sex
 Log(theta)
              -0.25032
                            0.21213
                                       -1.180 0.237979
Zero-inflation model coefficients (binomial with logit link):
 Estimate Std. Error z value Pr(>|z|)
(Intercept) -24.0614 317.6901 -0.076 0.940
                             0.7880
                                                   0.534
 Swimmer
                -0.4899
                                       -0.622
 Location
                11.8002
                           158.8409
                                        0.074
                                                   0.941
                0.1043
                              0.0859
                                        1.214
                                                   0.225
 Aae
sex
                -1.4259
                             1.3285
                                       -1.073
                                                  0.283
 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Theta = 0.7786
Number of iterations in BFGS optimization: 47 Log-likelihood: -440.9 on 11 Df
call:
glm(formula = Infections ~ Swimmer + Location + Age + Sex, family = poisson,
     data = earinfection)
Deviance Residuals:
 Min 1Q Median
-2.2645 -1.5740 -1.1165
                      Median
                                 3Q
0.5223
                                               Max
                                            6.5151
 Coefficients:
              (Intercept)
 Swimmer
                                       -4.670 3.01e-06 ***
 Location
              -0.48960
                            0.10483
               -0.02606
                             0.01222
                                       -2.132
                                                   0.033
Age
               0.02938
                            0.10916
                                       0.269
                                                   0.788
Sex
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
 (Dispersion parameter for poisson family taken to be 1)
Null deviance: 824.51 on 286
Residual deviance: 760.01 on 282
                                         degrees of freedom
                                         degrees of freedom
AIC: 1142.4
Number of Fisher Scoring iterations: 6
```