

SOCY7704: Regression Models for Categorical Data
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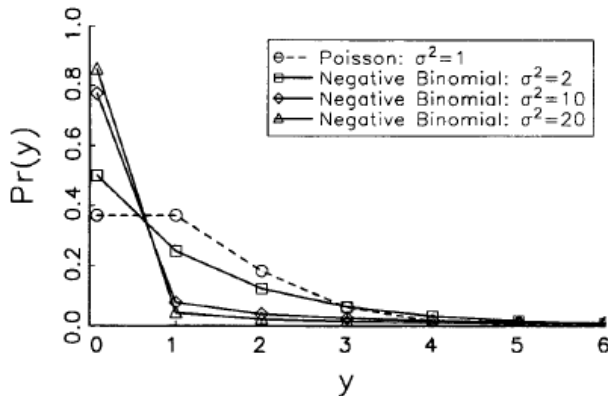
Count Data Models

Negative Binomial Model

Using Poisson, we attempted to account for some sources of heterogeneity – but the model doesn't fit very well. Maybe we didn't take into account all sources of heterogeneity – could try additional variables. That's important to explore, but rarely helps. In practice, Poisson regression model rarely fits due to overdispersion. One key process that often creates overdispersion is known as contagion – violation of the assumption of the independence of events. This assumption is often unrealistic; e.g. if you have your first child, that increases your chances of having your second.

To better model overdispersion from this and other sources, we can use negative binomial model. It allows taking into account unobserved heterogeneity. To do so, it introduces an additional parameter – alpha, known as the dispersion parameter. Increasing alpha increases the conditional variance of our count variable. If alpha is zero, the model becomes regular Poisson model. Here's a comparison of Poisson and negative binomial distributions with different variances for mean count=1 and mean count=10:

Panel A: $E(y)=1$



Panel B: $E(y)=10$

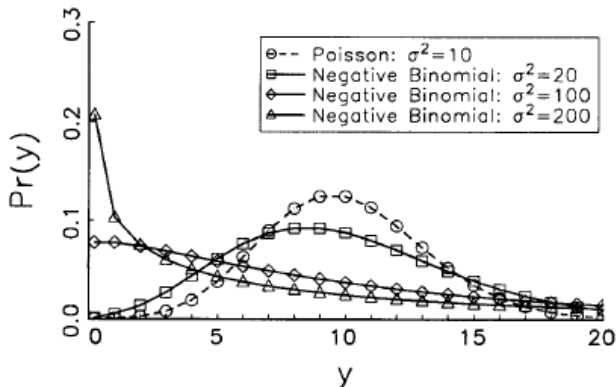
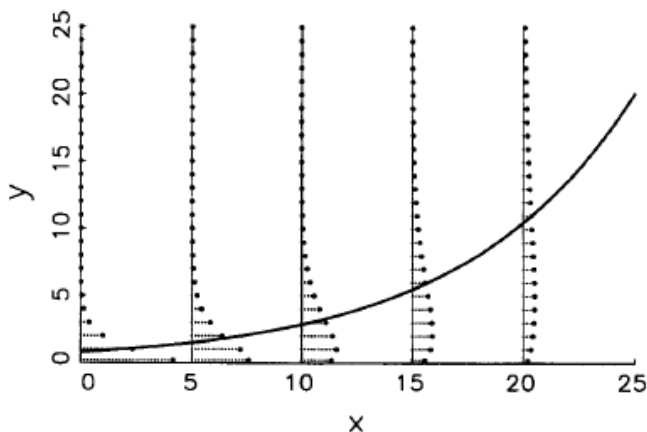


Figure 8.6. Comparisons of the Negative Binomial and Poisson Distributions

And here's an example of regression curves for negative binomial models:

Panel A: NBRM with $\alpha=0.5$



Panel B: NBRM with $\alpha=1.0$

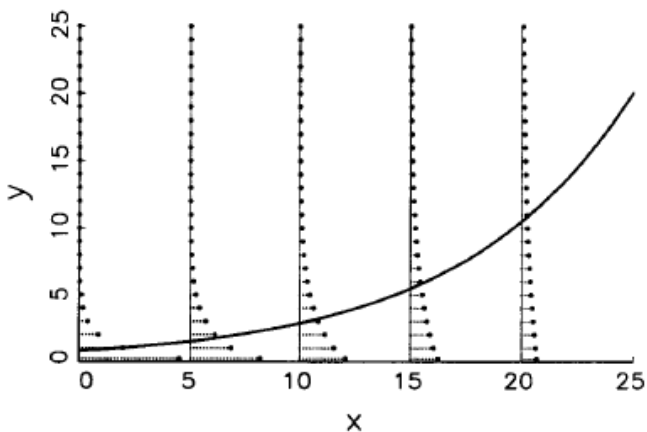


Figure 8.7. Distribution of Counts for the Negative Binomial Regression Model

Now let's run NB model for our data:

```
. nbreg childs sex married sibs born educ
Fitting Poisson model:
Iteration 0: log likelihood = -4784.5123
Iteration 1: log likelihood = -4784.5079
Iteration 2: log likelihood = -4784.5079
Fitting constant-only model:
Iteration 0: log likelihood = -5023.5027
Iteration 1: log likelihood = -4901.9594
Iteration 2: log likelihood = -4901.9154
Iteration 3: log likelihood = -4901.9154
Fitting full model:
Iteration 0: log likelihood = -4732.0308
Iteration 1: log likelihood = -4712.421
Iteration 2: log likelihood = -4711.6797
Iteration 3: log likelihood = -4711.6789
Iteration 4: log likelihood = -4711.6789
```

Negative binomial regression	Number of obs	=	2745
	LR chi2(5)	=	380.47
Dispersion = mean	Prob > chi2	=	0.0000
Log likelihood = -4711.6789	Pseudo R2	=	0.0388

childs	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
sex	.2086278	.0346569	6.02	0.000	.1407014	.2765542
married	.471206	.034682	13.59	0.000	.4032305	.5391816
sibs	.0397041	.0054244	7.32	0.000	.0290725	.0503358
born	-.2231164	.0616061	-3.62	0.000	-.3438622	-.1023706
educ	-.0616831	.0058316	-10.58	0.000	-.0731129	-.0502534
_cons	.9198597	.1211683	7.59	0.000	.6823743	1.157345
/lnalpha	-1.523939	.1086487			-1.736886	-1.310991
alpha	.2178522	.0236694			.1760678	.2695528

Likelihood-ratio test of alpha=0: chibar2(01) = 145.66 Prob>=chibar2 = 0.000

Or better yet, we will estimate this model with robust standard errors – it is recommended that we use them with negative binomial model in case the variance is misspecified.

```
. nbreg childs sex married sibs born educ, robust
Negative binomial regression          Number of obs   =       2745
Dispersion = mean                    Wald chi2(5)    =       386.44
Log pseudolikelihood = -4711.6789    Prob > chi2     =       0.0000
```

childs	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
sex	.2086278	.035025	5.96	0.000	.1399801	.2772755
married	.471206	.0348392	13.53	0.000	.4029225	.5394895
sibs	.0397041	.005216	7.61	0.000	.029481	.0499272
born	-.2231164	.0585515	-3.81	0.000	-.3378753	-.1083576
educ	-.0616831	.0060308	-10.23	0.000	-.0735032	-.049863
_cons	.9198597	.1225929	7.50	0.000	.6795821	1.160137
/lnalpha	-1.523939	.1167233			-1.752712	-1.295165
alpha	.2178522	.0254284			.1733033	.2738526

Interpretation of the results for negative binomial model is exactly the same as for Poisson model. But we have an extra line of output to interpret – the likelihood-ratio test. This allows us to see whether NB model should be used in place of regular Poisson. If probability is below the cutoff, it means that there is overdispersion (Alpha is not zero) and we should be using NB model rather than Poisson. Let's compare the coefficients to Poisson:

```
. est store nbreg
. qui poisson childs sex married sibs born educ
. est store poisson
. est table poisson nbreg, star b(%4.3f)
```

Variable	poisson	nbreg
childs		
sex	0.195***	0.209***
married	0.449***	0.471***
sibs	0.039***	0.040***
born	-0.221***	-0.223***
educ	-0.062***	-0.062***
_cons	0.955***	0.920***
lnalpha		
_cons		-1.524***

legend: * p<0.05; ** p<0.01; *** p<0.001

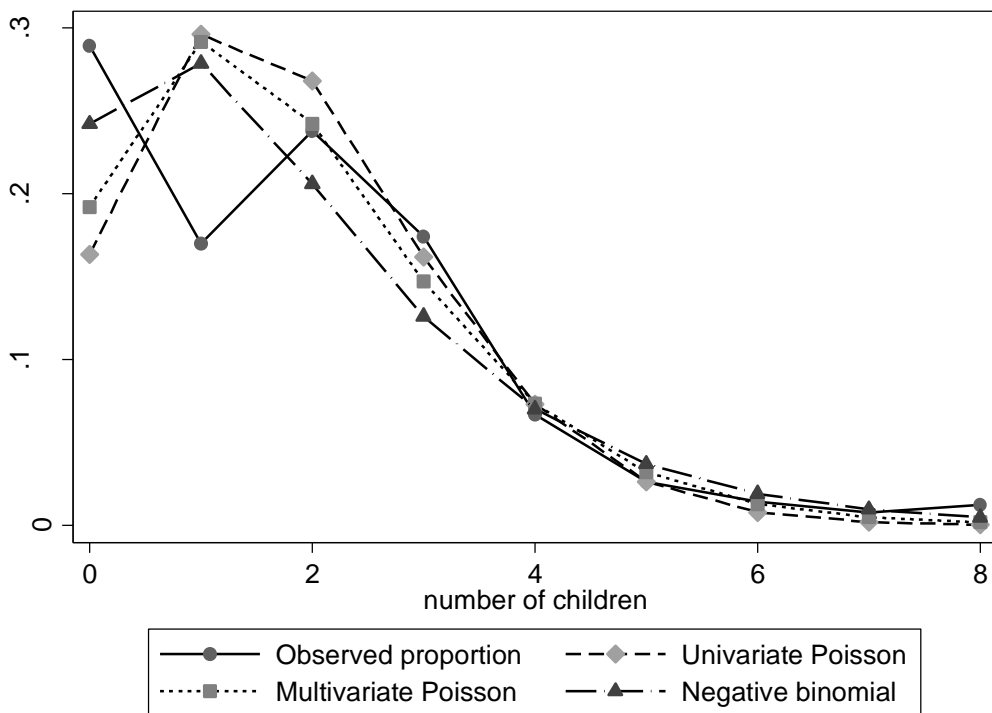
Now let's compare their performance graphically:

```
. mgen, pr(0/8) meanpred stub(nb_)
```

Predictions from:

Variable	Obs	Unique	Mean	Min	Max	Label
nb_val	9	9	4	0	8	number of children
nb_obeq	9	9	.1111111	.0080146	.2892532	Observed proportion
nb_oble	9	9	.7987047	.2892532	1	Observed cum. proportion
nb_preq	9	9	.1105054	.0049814	.2786995	Avg predicted Pr(y=#)
nb_prle	9	9	.7990764	.2423203	.9945486	Avg predicted cum. Pr(y=#)
nb_ob_pr	9	9	.0006057	-.108572	.0479451	Observed - Avg Pr(y=#)

```
. lab var nb_preq "Negative binomial"
. graph twoway connected poi_obeq poi_preq mpoi_preq nb_preq poi_val, ylabel(0 (.1) .3)
yttitle("Probability of Count")
```



The graph confirms the results of the alpha significance test: NB model does better than regular multivariate Poisson, especially with regard to dealing with 0s. But it still underpredicts zeros and overpredicts ones, and it underpredicts 2s and 3s (while Poisson was more on target).

Unfortunately, the goodness of fit tests that are available after Poisson are not available after negative binomial. But the significance test for alpha tells us if negative binomial model performs better than Poisson. We can also compare them using BIC:

```
. qui poisson child sex married sibs born educ
. qui fitstat, save
. qui nbreg child sex married sibs born educ
. fitstat, diff
```

	Current	Saved	Difference
Log-likelihood			
Model	-4711.679	-4784.508	72.829
Intercept-only	-4901.915	-5070.839	168.924

Chi-square				
D (df=2738/2739/-1)	9423.358	9569.016	-145.658	
Wald (df=5/5/0)	386.441	.	.	
p-value	0.000	0.000	.	

R2				
McFadden	0.039	0.056	-0.018	
McFadden (adjusted)	0.037	0.055	-0.018	
Cox-Snell/ML	0.129	0.188	-0.059	
Cragg-Uhler/Nagelkerke	0.133	0.193	-0.060	

IC				
AIC	9437.358	9581.016	-143.658	
AIC divided by N	3.438	3.490	-0.052	
BIC (df=7/6/1)	9478.781	9616.521	-137.740	

Note: Some measures based on pseudolikelihoods.

Difference of 137.740 in BIC provides very strong support for current model.

The interpretation tools for nbreg are the same as for Poisson; we can get IRR and use mtable, mchange, and mgen commands. We could also estimate this model with exposure.

As for diagnostics, everything is similar to Poisson, except for boxtid which doesn't work with nbreg. To obtain a GLM negative binomial model that's identical to the one estimated to nbreg, you need to specify the exact alpha to use – otherwise it uses the default value of 1 and the results differ. So here we use:

```
. glm childsex married sibs born educ, family(nb .2178552)
```

```
Generalized linear models          No. of obs      =      2745
Optimization      : ML              Residual df    =      2739
                                          Scale parameter =          1
Deviance          = 3284.463783      (1/df) Deviance = 1.199147
Pearson           = 2908.984543      (1/df) Pearson  = 1.062061

Variance function: V(u) = u+(.2178552)u^2      [Neg. Binomial]
Link function     : g(u) = ln(u)              [Log]
                                          AIC            = 3.437289
Log likelihood    = -4711.678905              BIC            = -18401.67
```

childsex	Coef.	OIM Std. Err.	z	P> z	[95% Conf. Interval]	

sex	.2086279	.0346384	6.02	0.000	.1407379	.2765179
married	.4712062	.0346364	13.60	0.000	.4033201	.5390924
sibs	.0397041	.0054238	7.32	0.000	.0290737	.0503346
born	-.2231165	.0616059	-3.62	0.000	-.3438618	-.1023712
educ	-.0616831	.0058316	-10.58	0.000	-.0731129	-.0502533
_cons	.9198593	.1211388	7.59	0.000	.6824317	1.157287

We can obtain residuals etc. after this.

In addition to regular nbreg where overdispersion is assumed to be constant, we can also use generalized negative binomial regression to model overdispersion (i.e., allow for different degree of overdispersion for different groups):

```
. gnbreg childsex married sibs born educ, lnalpha(sex married sibs born educ)
Generalized negative binomial regression      Number of obs  =      2745
                                          LR chi2(5)    =      222.46
                                          Prob > chi2   =      0.0000
```

Log likelihood = -4587.1261

Pseudo R2 = 0.0237

	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	

childs						
sex	.079685	.0354711	2.25	0.025	.0101628	.1492071
married	.3413691	.0387924	8.80	0.000	.2653374	.4174008
sibs	.0369471	.0047258	7.82	0.000	.0276847	.0462095
born	-.1967968	.0582151	-3.38	0.001	-.3108963	-.0826973
educ	-.0514978	.0056236	-9.16	0.000	-.0625199	-.0404758
_cons	1.085011	.1189463	9.12	0.000	.8518807	1.318142

lnalpha						
sex	-1.557369	.1884906	-8.26	0.000	-1.926804	-1.187934
married	-4.256861	.819715	-5.19	0.000	-5.863473	-2.650249
sibs	-.1051836	.0405024	-2.60	0.009	-.1845669	-.0258003
born	.1353893	.3910783	0.35	0.729	-.63111	.9018887
educ	.1619184	.0358938	4.51	0.000	.0915678	.232269
_cons	.3279141	.7155448	0.46	0.647	-1.074528	1.730356

Looks like overdispersion parameter varies by sex, marital status, number of siblings, and education, so the contagion process operates differently for different people (it is especially pronounced for men, unmarried people, those with fewer siblings, and those with more education).

Zero-Inflated Count Data Models

The problem that our negative binomial model still has – underpredicting zeros, overpredicting ones -- is very common and sometimes this problem can be very severe when there are a lot of zeros in the distribution. We can use zero-inflated count models to correct for that – they model two different processes. They assume two latent groups – one is capable of having positive counts, the other one is not – it will always have zero count. For example, some will have children eventually, but others do not have kids and cannot have them anymore or do not want to, so their count will always remain zero. But these two groups are latent – no information on their fertility situation or preferences. We can also have zeros in the first group. We can distinguish structural zeros (this behavior is not in this person’s repertoire at all) vs chance zeros (this behavior is in this person’s repertoire, but did not occur during the specified period). E.g.: “How many times last week did you smoke marijuana?” Some zeros mean the person never smokes it; other zeros mean the person does smoke but did not smoke last week.

Therefore, this model is a two-step process – first, you have to predict the membership in two groups – “always zero” and “not always zero” -- and second, predict the count in the “not always zero” group.

```
. zip childs sex married sibs born educ, inflate(sex married sibs born educ)
```

```
Zero-inflated poisson regression          Number of obs   =      2745
                                           Nonzero obs     =      1951
                                           Zero obs        =       794
```

```
Inflation model = logit                  LR chi2(5)      =      130.65
Log likelihood = -4524.192                Prob > chi2     =       0.0000
```

	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	

childs						
sex	.0014908	.0320997	0.05	0.963	-.0614234	.064405

married		.0307475	.0333411	0.92	0.356	-.0345999	.0960949
sibs		.0292838	.0045691	6.41	0.000	.0203286	.038239
born		-.1728303	.0563097	-3.07	0.002	-.2831953	-.0624654
educ		-.0382489	.0052824	-7.24	0.000	-.0486021	-.0278956
_cons		1.363043	.1094042	12.46	0.000	1.148615	1.577472

inflate							
sex		-1.267402	.1427508	-8.88	0.000	-1.547189	-.987616
married		-3.867796	.6722317	-5.75	0.000	-5.185346	-2.550246
sibs		-.0907598	.0284525	-3.19	0.001	-.1465256	-.034994
born		.3182067	.2733966	1.16	0.244	-.2176408	.8540542
educ		.1671403	.0267744	6.24	0.000	.1146635	.2196171
_cons		-.9103566	.5168716	-1.76	0.078	-1.923406	.102693

Note the inflate option we specified – we have to specify that option, it tells Stata what variables to use to predict the membership in “Always Zero” group. In this case, we used the same variables but we could have used a smaller subset of the variables or even different variables altogether.

We’ll return to interpreting this output. But let’s prepare to graphically examine the fit:

```
. mgen, pr(0/8) meanpred stub(zip_)
```

Predictions from:

Variable	Obs	Unique	Mean	Min	Max	Label
zip_val	9	9	4	0	8	number of children
zip_obeq	9	9	.1111111	.0080146	.2892532	Observed proportion
zip_oble	9	9	.7987047	.2892532	1	Observed cum. proportion
zip_preq	9	9	.1109995	.0021302	.2880608	Avg predicted Pr(y=#)
zip_prle	9	9	.7987461	.2880608	.9989958	Avg predicted cum. Pr(y=#)
zip_ob_pr	9	9	.0001116	-.021445	.0296168	Observed - Avg Pr(y=#)

```
. lab var zip_preq "ZIP"
```

We will also estimate a zero-inflated negative binomial model and then compare all of them.

```
. zinb childs sex married sibs born educ, inflate(sex married sibs born educ)
Zero-inflated negative binomial regression      Number of obs   =      2745
                                                Nonzero obs     =      1951
                                                Zero obs        =       794
Inflation model = logit                      LR chi2(5)      =      124.23
Log likelihood = -4522.91                    Prob > chi2     =       0.0000
```

childs		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
childs							
sex		.0060583	.0331917	0.18	0.855	-.0589961 .0711128	
married		.0346028	.0344018	1.01	0.314	-.0328234 .102029	
sibs		.0297016	.004743	6.26	0.000	.0204055 .0389977	
born		-.1730859	.0572733	-3.02	0.003	-.2853394 -.0608324	
educ		-.0384851	.0054302	-7.09	0.000	-.0491281 -.0278422	
_cons		1.347192	.1125643	11.97	0.000	1.12657 1.567814	

inflate							
sex		-1.290154	.1468538	-8.79	0.000	-1.577982 -1.002326	
married		-4.405718	1.215488	-3.62	0.000	-6.78803 -2.023406	
sibs		-.0911606	.02947	-3.09	0.002	-.1489207 -.0334006	
born		.3417874	.2818703	1.21	0.225	-.2106681 .894243	
educ		.1715742	.0277136	6.19	0.000	.1172565 .2258919	
_cons		-.9919407	.5360101	-1.85	0.064	-2.042501 .0586197	

/lnalpha		-3.718083	.6593754	-5.64	0.000	-5.010435 -2.425731	

alpha		.0242805	.0160099			.006668 .0884134	

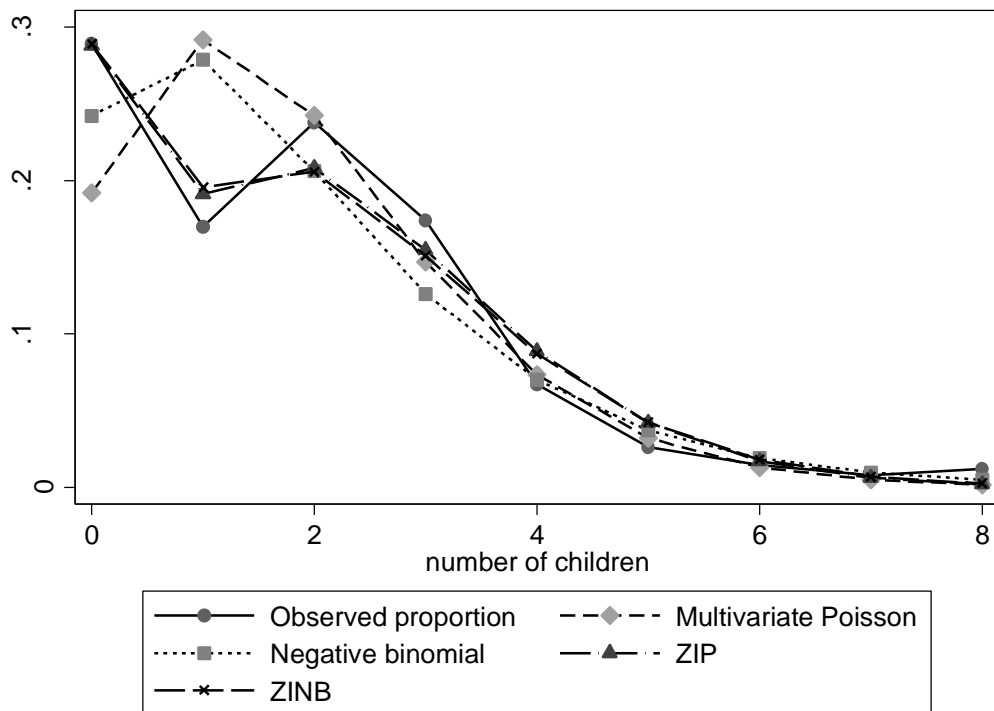
```
. mgen, pr(0/8) meanpred stub(zinb_)
```

```
Predictions from:
```

Variable	Obs	Unique	Mean	Min	Max	Label
zinb_val	9	9	4	0	8	number of children
zinb_obeq	9	9	.1111111	.0080146	.2892532	Observed proportion
zinb_oble	9	9	.7987047	.2892532	1	Observed cum. proportion
zinb_preq	9	9	.1109602	.0025516	.288929	Avg predicted Pr(y=#)
zinb_prle	9	9	.798788	.288929	.9986414	Avg predicted cum. Pr(y=#)
zinb_ob_pr	9	9	.000151	-.0256162	.0320836	Observed - Avg Pr(y=#)

```
. lab var zinb_preq "ZINB"
```

```
. graph twoway connected poi_obeq mpoi_preq nb_preq zip_preq zinb_preq poi_val, ylabel(0 (.1) .3) ytitle("Probability of Count")
```



Both ZIP and ZINB approximate the observed distribution much better than regular Poisson and NB models. We could also plot deviations from observed counts rather than actual counts and get comparisons of fit:

```
. countfit childs sex married sibs born educ, inflate(sex married sibs born educ)
```

Variable	PRM	NBRM	ZIP	ZINB
childs				
respondents sex	1.216	1.232	1.001	1.006
married	6.73	6.02	0.05	0.18
number of brothers and sisters	1.566	1.602	1.031	1.035
was r born in this country	15.54	13.59	0.92	1.01
highest year of school compl~d	1.039	1.041	1.030	1.030
born educ	9.14	7.32	6.41	6.26
sex	0.802	0.800	0.841	0.841
married	-4.23	-3.62	-3.07	-3.02
sibs	0.940	0.940	0.962	0.962
born educ	-12.81	-10.58	-7.24	-7.09

	Constant	2.598 9.45	2.509 7.59	3.908 12.46	3.847 11.97

lnalpha	Constant		0.218 -14.03		0.024 -5.64

inflate	respondents sex			0.282 -8.88	0.275 -8.79
	married			0.021 -5.75	0.012 -3.62
number of brothers and sisters				0.913 -3.19	0.913 -3.09
was r born in this country				1.375 1.16	1.407 1.21
highest year of school compl~d				1.182 6.24	1.187 6.19
	Constant			0.402 -1.76	0.371 -1.85

Statistics					
	alpha		0.218		
	N	2745	2745	2745	2745
	ll	-4784.508	-4711.679	-4524.192	-4522.910
	bic	9616.521	9478.781	9143.394	9148.749
	aic	9581.016	9437.358	9072.383	9071.821

Legend: b/t

Comparison of Mean Observed and Predicted Count

Model	Maximum Difference	At Value	Mean Diff
PRM	-0.122	1	0.028
NBRM	-0.109	1	0.027
ZIP	0.030	2	0.012
ZINB	0.032	2	0.013

PRM: Predicted and actual probabilities

Count	Actual	Predicted	Diff	Pearson
0	0.289	0.192	0.097	135.055
1	0.170	0.292	0.122	139.312
2	0.238	0.242	0.005	0.231
3	0.174	0.147	0.027	13.674
4	0.067	0.073	0.006	1.361
5	0.026	0.032	0.006	3.069
6	0.015	0.013	0.002	0.526
7	0.008	0.005	0.003	5.097
8	0.012	0.002	0.011	163.156
9	0.000	0.001	0.001	1.924
Sum	1.000	1.000	0.278	463.405

NBRM: Predicted and actual probabilities

Count	Actual	Predicted	Diff	Pearson
0	0.289	0.242	0.047	24.952
1	0.170	0.279	0.109	116.103
2	0.238	0.206	0.032	13.512
3	0.174	0.126	0.048	50.004
4	0.067	0.070	0.003	0.315

5	0.026	0.037	0.011	8.820
6	0.015	0.019	0.005	3.010
7	0.008	0.010	0.002	0.867
8	0.012	0.005	0.007	30.214
9	0.000	0.003	0.003	7.016

Sum	1.000	0.997	0.265	254.813
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ZIP: Predicted and actual probabilities

Count	Actual	Predicted	Diff	Pearson
0	0.289	0.288	0.001	0.014
1	0.170	0.191	0.021	6.403
2	0.238	0.208	0.030	11.561
3	0.174	0.155	0.019	6.512
4	0.067	0.089	0.021	14.210
5	0.026	0.042	0.016	16.286
6	0.015	0.017	0.003	1.083
7	0.008	0.006	0.002	1.298
8	0.012	0.002	0.010	135.546
9	0.000	0.001	0.001	1.886

Sum	1.000	1.000	0.124	194.798
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ZINB: Predicted and actual probabilities

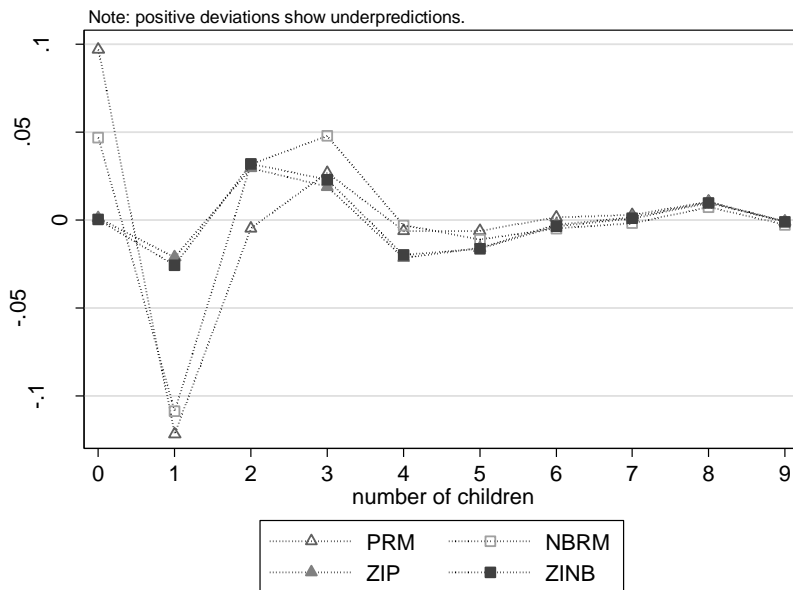
Count	Actual	Predicted	Diff	Pearson
0	0.289	0.289	0.000	0.001
1	0.170	0.196	0.026	9.202
2	0.238	0.206	0.032	13.730
3	0.174	0.151	0.023	9.695
4	0.067	0.087	0.020	12.320
5	0.026	0.042	0.016	16.787
6	0.015	0.018	0.003	1.855
7	0.008	0.007	0.001	0.389
8	0.012	0.003	0.010	104.052
9	0.000	0.001	0.001	2.445

Sum	1.000	1.000	0.132	170.477
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Tests and Fit Statistics

PRM	BIC=	9616.521	AIC=	9581.016	Prefer	Over	Evidence
vs NBRM	BIC=	9478.781	dif=	137.740	NBRM	PRM	Very strong
	AIC=	9437.358	dif=	143.658	NBRM	PRM	
	LRX2=	145.658	prob=	0.000	NBRM	PRM	p=0.000
vs ZIP	BIC=	9143.394	dif=	473.127	ZIP	PRM	Very strong
	AIC=	9072.383	dif=	508.632	ZIP	PRM	
	Vuong=	11.165	prob=	0.000	ZIP	PRM	p=0.000
vs ZINB	BIC=	9148.749	dif=	467.772	ZINB	PRM	Very strong
	AIC=	9071.821	dif=	509.195	ZINB	PRM	
NBRM	BIC=	9478.781	AIC=	9437.358	Prefer	Over	Evidence
vs ZIP	BIC=	9143.394	dif=	335.387	ZIP	NBRM	Very strong
	AIC=	9072.383	dif=	364.974	ZIP	NBRM	
vs ZINB	BIC=	9148.749	dif=	330.032	ZINB	NBRM	Very strong
	AIC=	9071.821	dif=	365.537	ZINB	NBRM	
	Vuong=	10.441	prob=	0.000	ZINB	NBRM	p=0.000

ZIP	BIC= 9143.394	AIC= 9072.383	Prefer	Over	Evidence
vs ZINB	BIC= 9148.749	dif= -5.355	ZIP	ZINB	Positive
	AIC= 9071.821	dif= 0.563	ZINB	ZIP	
	LRX2= 2.563	prob= 0.055	ZINB	ZIP	p=0.000



So now let's interpret this final model:

```
. zip childs sex married sibs born educ, inflate(sex married sibs born educ)
Zero-inflated poisson regression      Number of obs =      2745
                                       Nonzero obs   =      1951
                                       Zero obs       =       794
Inflation model = logit                LR chi2(5)      =     130.65
Log likelihood = -4524.192              Prob > chi2    =       0.0000
```

	childs	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]

childs						
	sex	.0014908	.0320997	0.05	0.963	-.0614234 .064405
	married	.0307475	.0333411	0.92	0.356	-.0345999 .0960949
	sibs	.0292838	.0045691	6.41	0.000	.0203286 .038239
	born	-.1728303	.0563097	-3.07	0.002	-.2831953 -.0624654
	educ	-.0382489	.0052824	-7.24	0.000	-.0486021 -.0278956
	_cons	1.363043	.1094042	12.46	0.000	1.148615 1.577472

inflate						
	sex	-1.267402	.1427508	-8.88	0.000	-1.547189 -.987616
	married	-3.867796	.6722317	-5.75	0.000	-5.185346 -2.550246
	sibs	-.0907598	.0284525	-3.19	0.001	-.1465256 -.034994
	born	.3182067	.2733966	1.16	0.244	-.2176408 .8540542
	educ	.1671403	.0267744	6.24	0.000	.1146635 .2196171
	_cons	-.9103566	.5168716	-1.76	0.078	-1.923406 .102693

The first set of coefficients is from the equation predicting counts for the “Not Always Zero” group. These show that number of siblings increases number of children and being foreign born and having more education decreases it. These coefficients can be interpreted the same way as regular Poisson coefficients.

The second set of coefficients is from the equation that predicts membership in “Always Zero” group. These can be interpreted as logit coefficients. Note that they predict zeros – so their sign will usually be the opposite to that of the coefficients in the upper half of the output. These show that women are less likely than men to be in “Always zero” group, married are less likely than single people to be in it, those with more siblings are also less likely to be in it, and those with more education are more likely to be in “Always zero” group.

To be able to interpret the size of these effects, let’s use listcoef to see IRR (but irr option is also available for zip and zinb commands themselves):

```
. listcoef
zip (N=2745): Factor Change in Expected Count
Observed SD: 1.6887584
Count Equation: Factor Change in Expected Count for Those Not Always 0
```

childs	b	z	P> z	e^b	e^bStdX	SDofX
sex	0.00149	0.046	0.963	1.0015	1.0007	0.4970
married	0.03075	0.922	0.356	1.0312	1.0154	0.4985
sibs	0.02928	6.409	0.000	1.0297	1.0919	3.0008
born	-0.17283	-3.069	0.002	0.8413	0.9512	0.2893
educ	-0.03825	-7.241	0.000	0.9625	0.8925	2.9741

```
Binary Equation: Factor Change in Odds of Always 0
```

Always0	b	z	P> z	e^b	e^bStdX	SDofX
sex	-1.26740	-8.878	0.000	0.2816	0.5326	0.4970
married	-3.86780	-5.754	0.000	0.0209	0.1454	0.4985
sibs	-0.09076	-3.190	0.001	0.9132	0.7616	3.0008
born	0.31821	1.164	0.244	1.3747	1.0964	0.2893
educ	0.16714	6.243	0.000	1.1819	1.6439	2.9741

Or better yet with percentages:

```
. listcoef, percent
zip (N=2745): Percentage Change in Expected Count
Observed SD: 1.6887584
Count Equation: Percentage Change in Expected Count for Those Not Always 0
```

childs	b	z	P> z	%	%StdX	SDofX
sex	0.00149	0.046	0.963	0.1	0.1	0.4970
married	0.03075	0.922	0.356	3.1	1.5	0.4985
sibs	0.02928	6.409	0.000	3.0	9.2	3.0008
born	-0.17283	-3.069	0.002	-15.9	-4.9	0.2893
educ	-0.03825	-7.241	0.000	-3.8	-10.8	2.9741

```
Binary Equation: Factor Change in Odds of Always 0
```

Always0	b	z	P> z	%	%StdX	SDofX
sex	-1.26740	-8.878	0.000	-71.8	-46.7	0.4970
married	-3.86780	-5.754	0.000	-97.9	-85.5	0.4985
sibs	-0.09076	-3.190	0.001	-8.7	-23.8	3.0008
born	0.31821	1.164	0.244	37.5	9.6	0.2893
educ	0.16714	6.243	0.000	18.2	64.4	2.9741

Each additional sibling increases one's number of kids by 3%, each year of education decreases it by 3.8%, and being foreign born decreases it by 16%. At the same time, women's odds of having no kids (being in always zero group) are 71.8% lower than men's, and the odds for married to be in always zero group are 97.9% lower than for single people. Further, each additional sibling decreases one's odds of not having kids by 8.7%, and each additional year of education increases those odds by 18.2%.

Further, as for regular Poisson, we can interpret predicted rates, predicted probabilities of specific counts, and changes in both rates and probabilities using `mtable`, `mchange`, and `mgen`. Predicted rates for by born and sex for married people:

```
. zip childs i.sex i.married sibs i.born educ, inflate(i.sex i.married sibs i.born
educ)
. mtable, at(sex=(1 2) born=(1 2) married==1) atmeans stat(ci)
Expression: Predicted number of childs, predict()
-----+-----
      |      sex      born      mu      ll      ul
-----+-----
      1 |          1          1      2.215      2.102      2.328
      2 |          1          2      1.849      1.645      2.053
      3 |          2          1      2.253      2.142      2.364
      4 |          2          2      1.891      1.684      2.099
Specified values of covariates
      | married      sibs      educ
-----+-----
Current |          1          3.6          13.4
```

Changes in predicted rates as well as marginal effects:

```
. mchange, amount(all)
zip: Changes in mu | Number of obs = 2745
Expression: Predicted number of childs, predict()
-----+-----
      |      Change      p-value
-----+-----
sex
female vs male |          0.332          0.000
married
  1 vs 0 |          0.801          0.000
sibs
  0 to 1 |          0.068          0.000
    +1 |          0.076          0.000
    +SD |          0.235          0.000
    Range |          2.547          0.000
  Marginal |          0.075          0.000
born
no vs yes |         -0.361          0.000
educ
  0 to 1 |         -0.153          0.000
    +1 |         -0.108          0.000
    +SD |         -0.310          0.000
    Range |         -2.411          0.000
  Marginal |         -0.110          0.000

Average prediction
      1.812
```

We interpret these results the same way as for regular Poisson model. Discrete changes and marginal effects are particularly useful in zero-inflated models because they combine the two equations to calculate the overall impact of each variable on the expected count. I would

recommend presenting marginal effects (average ones or at means) along with two sets of exponentiated coefficients (IRR and OR) when reporting the results of zero-inflated models.

We can also examine predicted probabilities of counts:

```
. mtable, at(sex=(1 2) born=(1 2) married==1) atmeans pr(0/4)
Expression: Pr(childs), predict(pr())
```

	sex	born	none	one	two	three	four
1	1	1	0.123	0.230	0.261	0.197	0.111
2	1	2	0.174	0.275	0.262	0.166	0.079
3	2	1	0.109	0.233	0.265	0.200	0.113
4	2	2	0.156	0.281	0.268	0.170	0.081

Specified values of covariates

	married	sibs	educ
Current	1	3.6	13.4

And changes in probabilities of counts:

```
. mchange, amount(all) pr(0/4)
zip: Changes in PrAny0 | Number of obs = 2745
Expression: Pr(childs = any 0), predict(pr(0))
```

	0	1	2	3	4
sex					
female vs male	-0.135	0.038	0.040	0.029	0.016
p-value	0.000	0.000	0.000	0.000	0.000
married					
1 vs 0	-0.314	0.084	0.092	0.069	0.040
p-value	0.000	0.000	0.000	0.000	0.000
sibs					
0 to 1	-0.016	-0.003	0.003	0.006	0.005
p-value	0.000	0.046	0.006	0.000	0.000
+1	-0.014	-0.004	0.001	0.005	0.005
p-value	0.000	0.003	0.249	0.000	0.000
+SD	-0.042	-0.013	0.002	0.014	0.016
p-value	0.000	0.001	0.529	0.000	0.000
Range	-0.282	-0.145	-0.079	0.035	0.111
p-value	0.000	0.000	0.007	0.094	0.000
Marginal	-0.015	-0.004	0.001	0.005	0.005
p-value	0.000	0.005	0.160	0.000	0.000
born					
no vs yes	0.067	0.026	-0.007	-0.028	-0.027
p-value	0.014	0.100	0.444	0.001	0.000
educ					
0 to 1	0.009	0.009	0.009	0.003	-0.004
p-value	0.000	0.000	0.000	0.141	0.001
+1	0.024	0.003	-0.004	-0.008	-0.007
p-value	0.000	0.019	0.000	0.000	0.000
+SD	0.074	0.008	-0.013	-0.024	-0.021
p-value	0.000	0.066	0.000	0.000	0.000
Range	0.399	0.109	0.007	-0.100	-0.139
p-value	0.000	0.000	0.728	0.000	0.000
Marginal	0.024	0.004	-0.003	-0.008	-0.007
p-value	0.000	0.009	0.000	0.000	0.000

Average predictions

	0	1	2	3	4
Pr(y base)	0.288	0.191	0.208	0.155	0.089

We can also use `mgen` to make all kinds of graphs for predicted rates and probabilities of counts and changes in these, like we did for regular Poisson.

We can also adjust our final, best-fitting model to exposure time:

```
. zip childs sex married sibs born educ, inflate(sex married sibs born educ)
exposure(reprage)
(31 missing values generated)
```

```
Zero-inflated poisson regression      Number of obs   =      2734
                                       Nonzero obs     =      1946
                                       Zero obs        =       788
Inflation model = logit              LR chi2(5)      =     119.40
Log likelihood = -4334.455            Prob > chi2     =       0.0000
```

	childs	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	

childs							
	sex	.0673734	.0319959	2.11	0.035	.0046625	.1300842
	married	.0372361	.0329312	1.13	0.258	-.0273079	.10178
	sibs	.0213414	.004529	4.71	0.000	.0124647	.0302181
	born	-.099738	.0548672	-1.82	0.069	-.2072757	.0077996
	educ	-.04122	.0051174	-8.05	0.000	-.0512498	-.0311901
	_cons	-1.996286	.1081046	-18.47	0.000	-2.208167	-1.784405
	reprage	(exposure)					

inflate							
	sex	-1.258563	.1789565	-7.03	0.000	-1.609311	-.9078144
	married	-7.69451	37.75966	-0.20	0.839	-81.70207	66.31305
	sibs	-.0533748	.0340675	-1.57	0.117	-.1201459	.0133964
	born	.3318979	.3383992	0.98	0.327	-.3313523	.9951481
	educ	.1963433	.0342241	5.74	0.000	.1292652	.2634213
	_cons	-1.914812	.6732486	-2.84	0.004	-3.234355	-.5952693

Note that the model changed – marriage that seemed so important is no longer significant, and neither is foreign born status! Looks like the effects of those were just function of age. Gender, siblings, and education predict the count, and gender and education predict the membership in always zero group.

Let's use `fitstat` to see whether this model with exposure performs better than the model without:

```
. quietly fitstat, save
. quietly zip childs sex married sibs born educ if reprage~=., inflate(sex married sibs
born educ)
```

Note: Here we limit the model without exposure only to those who don't miss data on `reprage` variable.

```
. fitstat, diff
```

		Current	Saved	Difference

Log-likelihood				
	Model	-4509.577	-4334.455	-175.121
	Intercept-only	-4825.719	-4825.719	0.000

Chi-square				
	D (df=2722/2722/0)	9019.153	8668.911	350.243
	LR (df=10/10/0)	632.285	982.528	-350.243
	p-value	0.000	0.000	.

R2				

	McFadden		0.066	0.102	-0.036
	McFadden (adjusted)		0.063	0.099	-0.036
	Cox-Snell/ML		0.206	0.302	-0.095
	Cragg-Uhler/Nagelkerke		0.213	0.311	-0.098

IC					
	AIC		9043.153	8692.911	350.243
	AIC divided by N		3.308	3.180	0.128
	BIC (df=12/12/0)		9114.116	8763.873	350.243

Difference of 350.243 in BIC provides very strong support for saved model.

We can see very strong support for the model with exposure, so we would select it as our final one.

Diagnostics for zero-inflated models:

Unfortunately, many tests and work-around solutions that worked for nbreg and poisson don't work for zip and zinb. One big problem is that zip and zinb cannot be modeled using GLM. We can still test for multicollinearity and use robust option for robust SE, but linearity diagnostics and those used to identify outliers and leverage points are not available here. So the strategy to use is:

1. Do the diagnostics using regular poisson or nbreg and then see if suggested fixes (e.g., a transformation or omitted leverage points) appear to improve the corresponding zero-inflated model.
2. Generate a dichotomy for 0 vs non-zero, run logit for that, and do diagnostics for logit as well (that would approximate the "Always zero" equation of ZIP and ZINB, and it is possible, for example, for a nonlinear relationship to exist in predicting counts but not predicting zeroes, or other way around).

Zero-truncated models

Sometimes we have count data that have no zeros at all, because we only start accumulating data once at least one count was observed. For example, the length of hospital stay cannot be 0 because we only start observing counts once a person is admitted. In such cases, zero-truncated models, implemented by ztp and ztnb commands, are useful. E.g., say, we only have data on the number of children after the person has their first one:

```
. gen childs0=childs
(5 missing values generated)
. replace childs0=. if childs==0
(799 real changes made, 799 to missing)

. ztp childs0 sex married sibs born educ
Zero-truncated Poisson regression          Number of obs   =       1951
                                           LR chi2(5)      =       168.39
                                           Prob > chi2     =       0.0000
                                           Pseudo R2      =       0.0262

Log likelihood = -3129.8812
```

childs0	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
sex	.0050533	.0341538	0.15	0.882	-.061887 .0719936
married	.0439347	.0344268	1.28	0.202	-.0235405 .11141
sibs	.0283134	.0047432	5.97	0.000	.019017 .0376098
born	-.1934924	.0631899	-3.06	0.002	-.3173423 -.0696426
educ	-.0403873	.0055964	-7.22	0.000	-.0513561 -.0294186
_cons	1.406071	.1183233	11.88	0.000	1.174161 1.63798


```

. ztnb childs0 sex married sibs born educ
Zero-truncated negative binomial regression      Number of obs   =      1951
                                                LR chi2(5)      =      114.29
Dispersion      = mean                        Prob > chi2     =      0.0000
Log likelihood = -3128.9162                    Pseudo R2      =      0.0179

```

childs0	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
sex	.0043327	.0352032	0.12	0.902	-.0646644	.0733297
married	.0440371	.0354945	1.24	0.215	-.0255309	.1136051
sibs	.0285975	.0049392	5.79	0.000	.0189169	.0382781
born	-.1951289	.0649357	-3.00	0.003	-.3224005	-.0678573
educ	-.0403866	.0057732	-7.00	0.000	-.0517018	-.0290714
_cons	1.398945	.1221116	11.46	0.000	1.15961	1.638279

/lnalpha	-3.811634	.7616972			-5.304533	-2.318735

alpha	.022112	.0168427			.004969	.098398

```

Likelihood-ratio test of alpha=0:  chibar2(01) =      1.93 Prob>=chibar2 = 0.082

```

Note that the results of these models look very similar to those from the count equations of zero-inflated Poisson and zero-inflated NB models.