

Poisson Regression with Incidence Rates

Poisson regression is used to model a non-negative integer **count variables** as the outcome. As mentioned earlier, in Public Health and Epidemiology disease incidence is often studied in relation to exposure over time

- Many deleterious exposures, as well as natural factors such as aging, will have bearing on the event count and must be accounted for when, say, comparing groups.
- Thus, rather than denominator N for a sample, the relevant denominator is the sum of exposure time over all N individuals, known as *person-time*. Rather than proportions, we have rates per unit of time at risk.
- This approach also accommodates different lengths of at risk time that may naturally occur.

Incidence Rates and Ratios

For incidence rate data

- data come in form of events and pyrs, there is no other 'N'. Data may be aggregated already (table form) or listed as individual cases with follow-up time (in which case there will be an 'N')
- latter form is needed for continuous exposure variables

Summary table for an epidemiological study of some exposure

	Exposure		Total
	yes	no	
case counts	c_e	c_u	$c_e + c_u = C$
person-years	pyr_e	pyr_u	$pyr_e + pyr_u = PYR$
Incidence rate	$IR_e = c_e/pyr_e$	$IR_u = c_u/pyr_u$	$IR = C/PYR$

Incidence Rates and Ratios

The following are cardiovascular disease events in relation to two potential risk factors, age group and obesity (y/n). The data (from SPRM Ch 9):

```
. list, clean noobs
```

AgeGroup	Obesity	cvd	pyrs	agegrp	obstat
60{64	Obese	10	245	1	1
60{64	Notobese	12	640	1	0
65{69	Obese	34	365	2	1
65{69	Notobese	45	520	2	0
70{74	Obese	40	250	3	1
70{74	Notobese	44	490	3	0

If we consider the obesity factor only, and sum the data to fill the table, we have

	Obesity		
	yes	no	Total
CVD cases	84	101	185
person-years	860	1650	2510
Incidence rate	.0976744	.0612121	.0737052

Previously we saw the ratio of count proportions as a natural summary from the Poisson model, showing how counts increase on a multiplicative scale

The incidence rate ratio is defined similarly as

$$\frac{IR_e}{IR_u} = \frac{.0977}{.0612} = 1.5956$$

Indicating about a 1.6 fold excess risk for the obese group

The Incidence Rate Ratio

STATA has modules to perform these types of epidemiologic summaries/analyses functions. The *incidence rate* command:

```
. ir cvd obstat pyrs
```

	obstat		
	Exposed	Unexposed	Total
-----+-----+-----			
cvd	84	101	185
pyrs	860	1650	2510
-----+-----+-----			
Incidence rate	.0976744	.0612121	.0737052
	Point estimate		[95% Conf. Interval]
	-----+-----		
Inc. rate diff.	.0364623		.0124039 .0605207
Inc. rate ratio	1.595671		1.180226 2.15264 (exact)
Attr. frac. ex.	.3733045		.1527047 .535454 (exact)
Attr. frac. pop	.1695004		
	-----+-----		

Poisson Regression Model (and other GLMs) - a note on model fitting

- The Poisson model and other GLMs are fit via *maximum likelihood estimation*. Least squares is an MLE estimator, but only for linear regression
- the *Likelihood Function* $L = f(Y_1, Y_2, \dots, Y_n | \omega)$ is a key quantity. It is the joint distribution of all the observations (data fixed), expressed as a function of all the parameters (ω). The method finds the values of the parameters (including β s) that most likely gave rise to the data. This is done by taking derivatives of $\log(L)$ with respect to parameters, setting equal to zero and solving.
- After the fit, L is a quantity (computed by plugging in those parameters) that can be used to compare models, measure 'fit', etc. We usually work with the *log likelihood*, provided with every model run.

Poisson Regression Model for Incidence Rates

We can fit the model as in the count case. **An important difference is the inclusion of the 'exposure' variable to indicate the person-years**

```
. poisson cvd obstat, exposure(pyrs) irr
. .
Iteration 1:   log likelihood = -42.981143
```

```
Poisson regression                               Number of obs   =           6
                                                LR chi2(1)        =           9.79
                                                Prob > chi2       =          0.0018
Log likelihood = -42.981143                    Pseudo R2        =          0.1022
```

cvd	IRR	Std. Err.	z	P> z	[95% Conf. Interval]	
obstat	1.595671	.2356291	3.16	0.002	1.194671	2.13127
_cons	.0612121	.0060908	-28.07	0.000	.0503663	.0743935
ln(pyrs)	1	(exposure)				

Note: _cons estimates baseline incidence rate.

The non-obese incidence rate and multiplicative effect of obesity (i.e.,

the IRR) are produced.

cvd	IRR	Std. Err.	z	P> z	[95% Conf. Interval]	
obstat	1.595671	.2356291	3.16	0.002	1.194671	2.13127
_cons	.0612121	.0060908	-28.07	0.000	.0503663	.0743935
ln(pyrs)	1	(exposure)				

Note: _cons estimates baseline incidence rate.

Note that .0612 or 6.1/100 person-years is the incidence rate in the non-obese and $.06125 \cdot 1.5956 = .09767$ or 9.8/100 person-years is the rate in the obese group, reproducing the numbers in the earlier table.

Poisson Model for Incidence Rates - add an ordinal predictor

We can add the age group variable as an ordinal predictor. Examine CVD by age:

```
. tabstat cvd pyrs, by(AgeGroup) stat(sum)  
Summary statistics: sum by categories of: AgeGroup
```

AgeGroup	cvd	pyrs
-----+-----		
60{64	22	885
65{69	79	885
70{74	84	740
-----+-----		
Total	185	2510

```
. display 22/885  
.02485876
```

```
. display 79/885  
.08926554
```

```
. display 84/740  
.11351351
```

Poisson Model for Incidence Rates - add an ordinal predictor

```
. poisson cvd obstat agegrp, exposure(pyrs) irr
```

```
Iteration 0:   log likelihood = -21.297411
Iteration 1:   log likelihood = -21.297215
Iteration 2:   log likelihood = -21.297215
```

```
Poisson regression               Number of obs   =           6
                                LR chi2(2)        =          53.15
                                Prob > chi2        =          0.0000
Log likelihood = -21.297215      Pseudo R2       =          0.5551
```

cvd		IRR	Std. Err.	z	P> z	[95% Conf. Interval]	
-----+-----							
obstat		1.535171	.2267304	2.90	0.004	1.149323	2.050554
agegrp		1.871195	.1845068	6.35	0.000	1.542366	2.27013
_cons		.0162154	.0040498	-16.50	0.000	.009939	.0264555
ln(pyrs)		1	(exposure)				

Note: _cons estimates baseline incidence rate.

Model here is on IRR scale

Poisson Model for Incidence Rates - Meaning of Predictor Coefficients

$\exp(\text{coefficient})$ yield the relative increase/decrease (i.e the IRR) for change in exposure level

- a. $\text{cons} = .0162$ - estimated mean CVD rate for non-obese, agegrp = 1
- b. $\beta_{\text{agegrp}} = 1.87$ - Indicating a 1.87-fold increase in CVD rate per age group increase
- c. $\beta_{\text{obstat}} = 1.535$ - Indicating a 1.54-fold increase in CVD for obese vs. nonobese- similar to earlier

Poisson Regression - Categorical Predictor for Age

```
. poisson cvd obstat age2 age3, exposure(pyrs) irr
```

```
Iteration 0:  log likelihood = -17.107802
Iteration 1:  log likelihood = -17.083109
Iteration 2:  log likelihood = -17.083097
Iteration 3:  log likelihood = -17.083097
```

```
Poisson regression                                Number of obs   =           6
                                                    LR chi2(3)      =        61.58
                                                    Prob > chi2     =         0.0000
Log likelihood = -17.083097                      Pseudo R2       =         0.6432
```

cvd	IRR	Std. Err.	z	P> z	[95% Conf. Interval]		
-----+-----							
obstat	1.468678	.2180134	2.59	0.010	1.097924	1.964629	
age2	3.399674	.8229289	5.06	0.000	2.115409	5.463615	
age3	4.453633	1.067596	6.23	0.000	2.784004	7.124574	
_cons	.0220038	.0048363	-17.36	0.000	.0143025	.0338521	
ln(pyrs)	1	(exposure)					

Note: _cons estimates baseline incidence rate.

Incidence Rates with Individual Case Data

The CVD data was aggregated by groups (age x obesity) with event counts. We can also work with individual data, where each individual either does or does not have the event over some follow-up time

Ex/ Endometrial cancer among women receiving tamoxifen/placebo in a breast cancer clinical trial. Tamoxifen is associated with increased risk, but is uncommon relative the the studied group ($N > 2800$) and amount of follow-up time:

```
.. list id trt menstat age bmi endo, clean
```

		id	trt	menstat	age	bmi	endo
.							
371.	635	1		3	67	36.6	0
372.	640	1		1	35	20.1	0
373.	642	1		3	50	19.6	0
374.	644	2		3	65	25.3	1
375.	646	1		3	58	35.7	0
376.	649	2		3	59	30.1	0
.							

Incidence Rates with Individual Case Data

The data summarized via Incidence Rate analysis (*timefree* is the exposure time)

```
. ir endo trtx timefree
```

Incidence-rate comparison

	trtx			
	Exposed	Unexposed	Total	
-----+-----+-----				
endo	13	3	16	
timefree	90322.5	79946.1	170268.6	
-----+-----+-----				
Incidence rate	.0001439	.0000375	.000094	
	Point estimate		[95% Conf. Interval]	
	-----+-----			
Inc. rate diff.	.0001064		.0000174	.0001954
Inc. rate ratio	3.835513		1.053989	20.98384 (exact)
Attr. frac. ex.	.7392787		.0512237	.9523443 (exact)
Attr. frac. pop	.6006639			
	-----+-----			

Incidence Rates with Individual Case Data

Use Poisson model:

```
. poisson endo trtx, exposure(timefree)
Iteration 0:   log likelihood = -93.429861
Iteration 1:   log likelihood = -93.428424
Iteration 2:   log likelihood = -93.428424
```

Poisson regression	Number of obs	=	1,395
	LR chi2(1)	=	5.58
	Prob > chi2	=	0.0182
Log likelihood = -93.428424	Pseudo R2	=	0.0290

endo	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
trtx	1.344303	.6405126	2.10	0.036	.0889214	2.599685
_cons	-10.1905	.5773502	-17.65	0.000	-11.32208	-9.05891
ln(timefree)	1	(exposure)				

```
. display exp(1.344303)
3.8355123
```

Incidence Rates with Individual Case Data

Poisson approach allows incorporation of covariates:

```
. poisson endo trtx age bmi, exposure(timefree) ir
```

```
Iteration 0: log likelihood = -92.189405
```

```
Iteration 1: log likelihood = -92.187682
```

```
Iteration 2: log likelihood = -92.187682
```

Poisson regression	Number of obs	=	1,395
	LR chi2(3)	=	8.06
	Prob > chi2	=	0.0448
Log likelihood = -92.187682	Pseudo R2	=	0.0419

endo	IRR	Std. Err.	z	P> z	[95% Conf. Interval]	
trtx	3.715235	2.381092	2.05	0.041	1.057917	13.04731
age	1.032185	.0297447	1.10	0.272	.9755024	1.092161
bmi	1.040999	.0452663	0.92	0.355	.9559539	1.133609
_cons	2.23e-06	4.38e-06	-6.62	0.000	4.71e-08	.0001052
ln(timefree)	1	(exposure)				

Note: _cons estimates baseline incidence rate.

Poisson Regression - Testing individual Coefficients

We can test the value of individual covariates as in SLR/MLR. We test

$$H_0 : \underset{\log}{IRR} = \underset{\log}{e^\beta} = 1$$

which is same as

$$H_0 : \log(IRR) = \beta = 0$$

vs.

$$H_A : \log(IRR) \neq \beta = 0$$

Theory from GLMs and the estimation method shows that β is \approx Normal. The test statistic (reported in the STATA/R output) for the above hypothesis is:

$$Z = \frac{\hat{\beta} - 0}{\text{se}(\hat{\beta})}$$

Poisson Regression - Testing the Whole Model

When we execute the model, the following summary precedes the

```
. poisson cvd obstat age2 age3, exposure(pyrs) irr
```

```
Iteration 0:    log likelihood = -17.107802
```

```
. . .
```

```
Iteration 3:    log likelihood = -17.083097
```

Poisson regression	Number of obs	=	6
	LR chi2(3)	=	61.58
	Prob > chi2	=	0.0000
Log likelihood = -17.083097	Pseudo R2	=	0.6432

The likelihood ratio test (LR above, equaling 61.58)) is a function of the difference in likelihoods for this model vs. a *null* model with no predictors. This is tantamount to the overall F test in LR.

To illustrate, we can run the two models and contrast

```
.* RUN null model
. poisson cvd, exposure(pyrs) irr
```

```
Iteration 0:   log likelihood = -47.874227
Iteration 1:   log likelihood = -47.874227
```

Poisson regression	Number of obs	=	6
	LR chi2(0)	=	-0.00
	Prob > chi2	=	.
Log likelihood = -47.874227	Pseudo R2	=	-0.0000

```
. * SAVE it
. est store nul
.
.* RUN full model
. poisson cvd obstat age2 age3, exposure(pyrs) irr
```

```
Iteration 0:   log likelihood = -17.107802
. . .
Iteration 3:   log likelihood = -17.083097
```

Poisson regression	Number of obs	=	6
	LR chi2(3)	=	61.58
	Prob > chi2	=	0.0000

```

Log likelihood = -17.083097                                Pseudo R2          =      0.6432
. . .
. * SAVE it
. est store big
.
.* CONTRAST these models
. lrtest big nul

Likelihood-ratio test                                LR chi2(3)  =      61.58
(Assumption: nul nested in big)                     Prob > chi2 =      0.0000

```

The above computation (test statistic is)

$$D = -2\{\log\hat{L}_n - \log\hat{L}_b\}$$

which here is

$$-2\{-47.874 - (-17.083)\} = 61.58$$

This is compared to a χ^2 statistic with degrees of freedom equal to the number of parameters (3 here). Result is to reject the null hypothesis that the likelihood ratio is 1.

Poisson Regression - Testing Subsets of Parameters

Contrasting different models follows the same strategy - contrasting the likelihood values between models.

- For example, to test the contribution of the categorical age variable (2 indicators), we can run the two models and compute as above.
- Alternatively, we can use a post-estimation test as done earlier in SLR/MLR. This is a different test (Wald test) but inference should be similar to LR test.

Poisson Regression - Testing Subsets of Parameters

```
. poisson cvd obstat age2 age3, exposure(pyrs) irr
```

```
. . .
```

```
Iteration 3:   log likelihood = -17.083097
```

Poisson regression	Number of obs	=	6
	LR chi2(3)	=	61.58
	Prob > chi2	=	0.0000
Log likelihood = -17.083097	Pseudo R2	=	0.6432

cvd	IRR	Std. Err.	z	P> z	[95% Conf. Interval]		
-----+-----							
obstat	1.468678	.2180134	2.59	0.010	1.097924	1.964629	
age2	3.399674	.8229289	5.06	0.000	2.115409	5.463615	
age3	4.453633	1.067596	6.23	0.000	2.784004	7.124574	
_cons	.0220038	.0048363	-17.36	0.000	.0143025	.0338521	
ln(pyrs)	1	(exposure)					

```
. test age2 age3
```

```
( 1)  [cvd]age2 = 0
```

```
( 2)  [cvd]age3 = 0
```

```

      chi2( 2) =    38.89
Prob > chi2 =    0.0000

```

Poisson Regression - Model and Coefficients

The LR test would use the same steps as earlier:

1. Run larger model (with age group vars and obesity), save log likelihood value (it is -17.083097)
2. Run model omitting age group vars, save likelihood value (it is -42.981143)
3. Compute $D = -2(-42.981143 - (-17.083097)) = 51.80$
4. Result is larger than χ^2 with 2 df, then age variable(s) are significantly contributing to model fit

Both the Wald and LR test conclude: keep age variables

Poisson Regression - Model Fit

To help assess the fit of the model, the `estat gof` command can be used to obtain the goodness-of-fit χ^2 test. This is **not** a test of the model coefficients, but rather a test of the model form: Does the Poisson model form fit our data? Thus, large p-value indicates good fit.

```
. estat gof
```

```
Deviance goodness-of-fit = 3.49824  
Prob > chi2(2)          = 0.1739
```

```
Pearson goodness-of-fit = 3.525841  
Prob > chi2(2)          = 0.1715
```

A statistically significant (small p-value) here would indicate that the data do not fit the model well. In that situation, we may try to determine if there are omitted predictor variables, if our linearity assumption holds and/or if the conditional mean and variance of outcome are very different.

Poisson Regression - Model Fit

To see the basis of the fit test, we can look at observed vs, predicted values:

```
. predict cvd_pred
(option n assumed; predicted number of events)
. list AgeGroup ObesityStatus cvd cvd_pred, noobs clean
```

AgeGroup	ObesityStatus	cvd	cvd_pred
60{64	Obese	10	7.91759
60{64	Notobese	12	14.0824
65{69	Obese	34	40.10115
65{69	Notobese	45	38.89889
70{74	Obese	40	35.98157
70{74	Notobese	44	48.01822

```
. gen chipart = (cvd - cvd_pred)^2 / (cvd_pred)
. tabstat chipart, stat(sum)
```

variable	sum
-----+-----	
chipart	3.52584

Note: This is the Pearson χ^2 sum that we compute for χ^2 tests in frequency tables

Fitting as a GLM

An alternative way to fit Poisson regression is using the “glm” function (Stata or R), specifying which “family” to use (default is linear regression).

```
. glm cvd obstat age2 age3, family(Poisson) exposure(pyrs)
```

```
Iteration 0:  log likelihood = -17.619973
Iteration 1:  log likelihood = -17.083841
Iteration 2:  log likelihood = -17.083097
Iteration 3:  log likelihood = -17.083097
```

Generalized linear models	Number of obs	=	6
Optimization : ML	Residual df	=	2
	Scale parameter	=	1
Deviance = 3.498240465	(1/df) Deviance	=	1.74912
Pearson = 3.525841336	(1/df) Pearson	=	1.762921

Variance function: $V(u) = u$	[Poisson]
Link function : $g(u) = \ln(u)$	[Log]

	AIC	=	7.027699
Log likelihood = -17.08309669	BIC	=	-.0852785

		OIM					
	cvd	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
obstat		.3843624	.1484419	2.59	0.010	.0934215	.6753032
age2		1.223679	.2420611	5.06	0.000	.7492484	1.698111
age3		1.49372	.2397136	6.23	0.000	1.02389	1.96355
_cons		-3.816539	.219792	-17.36	0.000	-4.247323	-3.385754
ln(pyrs)		1	(exposure)				

- Estimates are same as earlier. Again, β s are in log(rates) on an additive scale.
- The overall fit statistics as well as other measures are provided.

Fitting as a GLM

The model executed in R:

```
> library(foreign)
> cvd <- read.dta("CVD_factors.dta")
> cvd
```

	AgeGroup	ObesityStatus	cvd	pyrs	agegrp	obstat	age2	age3	_est_nul	_est_big	rate	cvd_pred
1	60{64	Obese	10	245	1	1	0	0	1	1	0.04081633	7.91759
2	60{64	Notobese	12	640	1	0	0	0	1	1	0.01875000	14.08240
3	65{69	Obese	34	365	2	1	1	0	1	1	0.09315068	40.10115
4	65{69	Notobese	45	520	2	0	1	0	1	1	0.08653846	38.89889
5	70{74	Obese	40	250	3	1	0	1	1	1	0.16000000	35.98157
6	70{74	Notobese	44	490	3	0	0	1	1	1	0.08979592	48.01822

```
      chipart  lnrate2
1 0.5476958 1.817766
2 0.3079306 1.857163
3 0.9282536 5.762812
4 0.9569315 7.195592
5 0.4487789 7.244460
6 0.3362491 7.103184
>
> Pois <- glm(cvd ~ offset(log(pyrs)) + obstat + age2 + age3, data=cvd, family=poisson)
>
```

```

>
> Pois

Call:  glm(formula = cvd ~ offset(log(pyrs)) + obstat + age2 + age3,
          family = poisson, data = cvd)

Coefficients:
(Intercept)      obstat      age2      age3
    -3.8165      0.3844      1.2237      1.4937

Degrees of Freedom: 5 Total (i.e. Null);  2 Residual
Null Deviance:      65.08
Residual Deviance:  3.498  AIC: 42.17

```

Poisson Regression - summary measures

In SLR/MLR, we have a partition of variability captured by R^2 . In Poisson regression (and other log-linear models), analogues to R^2 have been sought. One simple one is

$$R_{pseud}^2 = 1 - \frac{\log L(\hat{\beta})}{\log \hat{L}_0}$$

where $\log L(\hat{\beta})$ is the log likelihood for the current model and $\log \hat{L}_0$ is the log likelihood for the null model.

- Generally, these measures are not as reliable as fits measures as in the linear regression setting (although R^2 can be misleading there too)

Poisson Regression - summary measures

Ex/ for the full model (age groups and obesity vs. null model we have

$$R_{pseud}^2 = 1 - \frac{-17.083096}{-47.874227}$$

$$= 1 - 0.3568 = 0.6432$$

As given in the output for the model of interest:

```
. poisson cvd obstat age2 age3, exposure(pyrs) irr
. . .
```

Poisson regression	Number of obs	=	6
	LR chi2(3)	=	61.58
	Prob > chi2	=	0.0000
Log likelihood = -17.083097	Pseudo R2	=	0.6432
. . . .			

Alternate Models

Fit the Negative Binomial model (note: for this dist'n, variance increases as mean increases)

```
. nbreg cvd obstat age2 age3, exposure(pyrs)
```

Fitting Poisson model:

```
Iteration 0:  log likelihood = -17.107802
Iteration 1:  log likelihood = -17.083109
Iteration 2:  log likelihood = -17.083097
Iteration 3:  log likelihood = -17.083097
```

Fitting constant-only model:

```
Iteration 0:  log likelihood = -26.909909
Iteration 1:  log likelihood = -25.989194
Iteration 2:  log likelihood = -25.570977
Iteration 3:  log likelihood = -25.567453
Iteration 4:  log likelihood = -25.567453
```

Fitting full model:


```

Iteration 0:  log likelihood = -22.857883
Iteration 1:  log likelihood = -19.909726
Iteration 2:  log likelihood = -19.73484 (not concave)
Iteration 3:  log likelihood = -19.347292 (not concave)
Iteration 4:  log likelihood = -17.635909
Iteration 5:  log likelihood = -17.197221
Iteration 6:  log likelihood = -17.112039
Iteration 7:  log likelihood = -17.089736
Iteration 8:  log likelihood = -17.084648
Iteration 9:  log likelihood = -17.083448
Iteration 10: log likelihood = -17.083167
Iteration 11: log likelihood = -17.083109
Iteration 12: log likelihood = -17.083098
Iteration 13: log likelihood = -17.083096 (not concave)
Iteration 14: log likelihood = -17.083096

```

```

Negative binomial regression      Number of obs      =           6
                                LR chi2(3)                =          16.97
Dispersion      = mean          Prob > chi2          =          0.0007
Log likelihood = -17.083096     Pseudo R2           =          0.3318

```

```

-----+-----
            cvd |      Coef.   Std. Err.      z    P>|z|     [95 Conf. Interval]
-----+-----
      obstat |   .3843709   .148442    2.59   0.010   .0934299   .6753118

```

age2		1.223679	.2420612	5.06	0.000	.7492478	1.69811
age3		1.493717	.2397137	6.23	0.000	1.023887	1.963547
_cons		-3.816542	.2197921	-17.36	0.000	-4.247327	-3.385758
ln(pyrs)		1	(exposure)				
-----+-----							
/lnalpha		-18.60187	1339.439			-2643.855	2606.651
-----+-----							
alpha		8.34e-09	.0000112			0	.
-----+-----							
LR test of alpha=0: chibar2(01) = 6.8e-07				Prob >= chibar2 = 0.500			

Note: Poisson is a special case when $\alpha = 0$ - test above is for $H_0 : \alpha = 0$, which is NOT rejected. This means Poisson model is adequate. Parameters are very similar to Poisson model fit earlier.

Summary – Poisson Regression and GLMs

The Poisson data-based model provided a strong inferential and data exploratory tool for events in relation to person-time of exposure

When there seems to be an issue of bad fit, we should first check if our model is appropriately specified, such as omitted variables and functional form, and then consider variations on the model that may fit better

The Poisson provides a bridge between linear regression, discrete event counts, and rates of failure, which relates to time to event (survival) data

Summary – Courses Beyond Linear Regression

This course will briefly introduce variations on linear regression. Further development of these methods and others are extensively covered in other University of Chicago courses such as:

- *Categorical (Discrete) Data Analysis* - all types of discrete outcome variables - binomial, multinomial, ordinal, counts
- *Generalized Linear Models* - all GLMs
- *Biostatistical Methods* - logistic, Poisson, hazard (failure rate, time to event) in health science context
- *Applied Survival Analysis* - analysis of time to event data, including regression methods for these data
- *Applied Longitudinal Data Analysis* - extending regression to repeated measurements on Y