# WWS 509 Generalized Linear Models: Precept 4 Section 3.5

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#### 1 Introducing the Data

Today we will be looking at the same data as last week, women giving birth after HIV testing. For today, I have created group data, based on age groups (15-19, 20-24, 25-29, 30-34, 35-39, 40-44) and HIV status.

#### 2 Deviance

Deviance is the goodness of fit likelihood ratio chi-square statistic

- With a degree of freedom determined from the number of parameters added to the model
- $\bullet$  To see if a model fits, you want to look at the deviance and check the  $\chi^2$  distribution.

Regardless if the group or individual data, the estimated effects, the standard errors, and likelihood ratio test based on differences between deviances will be the same (but the deviances will change).

## 3 Categorical Age and Binary HIV status

There are 5 potential models from this data, what are they?

Model	Notation	$\operatorname{logit}(\pi_{ij})$	Deviance	D.F.

Using the above table:

- 1. How would I test the gross effect of age?
  - (a) Test if the model including only age fits the data.
- 2. How would I test the gross effect of HIV?
  - (a) Test if the model including only HIV fits the data.
- 3. Which variable, HIV or age, explains more (only looking at the single variable models)?
  - (a) Can you test this?

Now look at the additive model in the table:

- 1. What can this model tell you?
  - (a) What is the effect of adding age to the model containing HIV status?
  - (b) What is the effect of adding HIV to the model containing age?
- 2. Does this model fit the data?

Now finally, look at the saturated model in the table:

- 1. How much deviance is in this model?
- 2. Is this model useful for interpretation?

### 4 Linear Age and Binary HIV status

Now age as a categorical variable uses up many of our degrees of freedom, so lets see if we can model things a bit differently. Refer to the output in Appendix

B. Fill in the deviance table for all the models in Appendix B (and don't forget to include the null model from Appendix A).

Model	Notation	$\operatorname{logit}(\pi_{ij})$	Deviance	D.F.	

- 1. What do you think about modeling age linearly?
  - (a) Is it an improvement over the null?
  - (b) Does this model fit the data?

I added HIV into the model:

- 1. Is this model an improvement over the previous model?
  - (a) Does this model fit the data?

Lets now try an interaction term?

- 1. Is the interaction term significant (look at the output)?
  - (a) Does this model offer an improvement over the additive model?

OK, linear effect of age and childbearing, not a good idea. But this doesn't mean we have to return to our categorical variables. I add a quadratic term for age.

- 1. How did I create this term in Stata?
- 2. What do you think about this model?
  - (a) Does this model offer an improvement over the model of age as a linear predictor?

For the last regression, I model age, age squared, and HIV status.

- 1. Does this model offer an improvement over the model of age and age squared?
- 2. Does this model fit the data?

# **Appendices**

#### Appendix A: Grouped Factor Data Stata Output

```
. gen chi_dum=.
(3307 missing values generated)
. replace chi_dum=1 if child_born!=0
(1114 real changes made)
. replace chi_dum=0 if child_born==0
(2193 real changes made)
```

. tab age\_range, gen(age\_group)

age_range	1	Freq.	Percent	Cum.
	-+			
17.5	1	817	24.71	24.71
22.5	1	582	17.60	42.30
27.5	1	569	17.21	59.51
32.5	1	494	14.94	74.45
37.5	1	396	11.97	86.42
42.5	1	449	13.58	100.00
	+			
Total	1	3,307	100.00	

```
. gen n=1
```

. collapse (sum) chi\_dum n, by (age\_range hiv5)

```
. glm chi_dum, family(binomial n)
```

```
Iteration 0: log likelihood = -216.53224
Iteration 1: log likelihood = -212.84368
Iteration 2: log likelihood = -212.8429
Iteration 3: log likelihood = -212.8429
```

Generalized linear	models	No. of obs	=	12
Optimization :	ML	Residual df		11
		Scale parameter	=	1
Deviance =	366.1140405	(1/df) Deviance	=	33.28309
Pearson =	349.6392631	(1/df) Pearson	=	31.78539

Variance function: V(u) = u\*(1-u/n) [Binomial] Link function : g(u) = ln(u/(n-u)) [Logit]

Log likelihood = -212.8429045				AIC BIC		35.64048 338.7801
	   Coef.				[95% Conf.	Interval]
	6773133				7494246	605202
. glm chi_dum	hiv5, family	(binomial n)	)			
Iteration 0: Iteration 1: Iteration 2: Iteration 3:	log likelih log likelih log likelih log likelih	pod = -206.3 pod = -206.3	18839 18768			
Generalized 1: Optimization Deviance	: ML = 352.80			Residence Reside	of obs = dual df = e parameter = f) Deviance =	10 1 35.28036
Pearson  Variance function		u*(1-u/n)			f) Pearson = omial] it]	33.49689
Log likelihoo	d = -206.18	76824		AIC BIC		34.69795 327.9545
chi_dum	   Coef.	OIM Std. Err.	z	P> z	[95% Conf.	Interval]
	5553665  641307					

. tab age\_range, gen(age\_group)

age_range	Freq.	Percent	Cum.
17.5	2	16.67	16.67
22.5	2	16.67	33.33
27.5	2	16.67	50.00
32.5	2	16.67	66.67

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37.5 42.5	•	2	16.67 16.67	83.33 100.00
Total		12	100.00	

. drop age\_group1

. glm chi\_dum age\_group\*, family(binomial n)

Iteration 0: log likelihood = -49.074751
Iteration 1: log likelihood = -49.0383
Iteration 2: log likelihood = -49.0383

 Generalized linear models
 No. of obs = 12

 Optimization : ML
 Residual df = 6

 Scale parameter = 1

 Deviance = 38.50483154
 (1/df) Deviance = 6.417472

 Pearson = 36.28561409
 (1/df) Pearson = 6.047602

Variance function: V(u) = u\*(1-u/n) [Binomial] Link function : g(u) = ln(u/(n-u)) [Logit]

AIC = 9.17305 Log likelihood = -49.03830004 BIC = 23.59539

. glm chi\_dum age\_group\* hiv5, family(binomial n)

Iteration 0: log likelihood = -31.604792
Iteration 1: log likelihood = -31.585329
Iteration 2: log likelihood = -31.585328

Generalized linear models No. of obs = 12 Optimization : ML Residual df = 5 Scale parameter = 1

Deviance = 3.59888798 (1/df) Deviance = .7197776Pearson = 4.468746092 (1/df) Pearson = .8937492

Variance function: V(u) = u\*(1-u/n) [Binomial] Link function : g(u) = ln(u/(n-u)) [Logit]

AIC = 6.430888 Log likelihood = -31.58532826 BIC = -8.825645

   chi_dum 	Coef.	OIM Std. Err.	z	P> z	[95% Conf.	Interval]
age_group2	1.210116	.1264414	9.57	0.000	.9622954	1.457937
age_group3	1.705656	.1267314	13.46	0.000	1.457268	1.954045
age_group4	1.624498	.1312985	12.37	0.000	1.367158	1.881838
age_group5	1.201172	.1407213	8.54	0.000	.9253632	1.47698
age_group6	1244814	.1630386	-0.76	0.445	4440312	.1950684
hiv5	9173204	.1638805	-5.60	0.000	-1.23852	5961205
_cons	-1.585984	.0934129	-16.98	0.000	-1.76907	-1.402898

. gen age\_hiv2=age\_group2\*hiv5

. gen age\_hiv3=age\_group3\*hiv5

. gen age\_hiv4=age\_group4\*hiv5

. gen age\_hiv5=age\_group5\*hiv5

. gen age\_hiv6=age\_group6\*hiv5

. glm chi\_dum age\_group\* hiv5 age\_hiv\*, family(binomial n)

Iteration 0: log likelihood = -29.786201
Iteration 1: log likelihood = -29.785884
Iteration 2: log likelihood = -29.785884

 Generalized linear models
 No. of obs = 12

 Optimization : ML
 Residual df = 0

 Scale parameter = 1
 1

 Deviance = 1.69961e-14
 (1/df) Deviance = .

 Pearson = 1.01332e-16
 (1/df) Pearson = .

Variance function: V(u) = u\*(1-u/n) [Binomial] Link function : g(u) = ln(u/(n-u)) [Logit]

Log likelihood	= -29.785	88427		AIC BIC		0.001011
1		OIM				
chi_dum	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
age_group2	1.234313	.1278387	9.66	0.000	. 983754	1.484873
age_group3	1.730612	.1291826	13.40	0.000	1.477418	1.983805
age_group4	1.638989	.1343145	12.20	0.000	1.375737	1.90224
age_group5	1.182523	.1444718	8.19	0.000	.8993634	1.465682
age_group6	1099815	.1652238	-0.67	0.506	4338142	.2138511
hiv5	.3448405	.8072783	0.43	0.669	-1.237396	1.927077
age_hiv2	-1.630209	.94745	-1.72	0.085	-3.487177	.2267589
age_hiv3	-1.418832	.863965	-1.64	0.101	-3.112172	.2745082
age_hiv4	-1.290682	.8622822	-1.50	0.134	-2.980724	.3993601
age_hiv5	9483295	.8766014	-1.08	0.279	-2.666437	.7697778
age_hiv6	-1.345306	1.097037	-1.23	0.220	-3.495458	.8048466
_cons	-1.597603	.0940268	-16.99	0.000	-1.781893	-1.413314

# Appendix B: Grouped Data with Linear Age Stata Output

```
. glm chi_dum age_range, family(binomial n)
Iteration 0:
              log likelihood = -214.93171
Iteration 1:
              log likelihood = -210.95632
Iteration 2:
              log likelihood = -210.95541
Iteration 3:
              log likelihood = -210.95541
Generalized linear models
                                                 No. of obs
                                                                          12
Optimization : ML
                                                 Residual df
                                                                          10
                                                 Scale parameter =
                = 362.3390561
Deviance
                                                 (1/df) Deviance = 36.23391
                = 345.5239143
                                                 (1/df) Pearson = 34.55239
Pearson
Variance function: V(u) = u*(1-u/n)
                                                  [Binomial]
Link function : g(u) = \ln(u/(n-u))
                                                  [Logit]
```

AIC = 35.49257 Log likelihood = -210.9554123BIC = 337.49 OIM chi\_dum | Coef. Std. Err. z P>|z| [95% Conf. Interval] age\_range | .0082429 .0042403 1.94 0.052 -.000068 .0165538 \_cons | -.9100142 .125642 -7.24 0.000 -1.156268 -.6637604 . glm chi\_dum age\_range hiv5, family(binomial n) Iteration 0:  $log\ likelihood = -207.24585$ Iteration 1: log likelihood = -203.37119Iteration 2: log likelihood = -203.37042Iteration 3: log likelihood = -203.37042 No. of obs = Residual df = Generalized linear models 12 Optimization : ML 9 Scale parameter = 1 = 347.1690663 (1/df) Deviance = 38.57434 Deviance = 328.9774173 Pearson (1/df) Pearson = 36.55305 Variance function: V(u) = u\*(1-u/n)[Binomial] Link function :  $g(u) = \ln(u/(n-u))$ [Logit] AIC = 34.39507 Log likelihood = -203.3704174BIC = 324.8049 \_\_\_\_\_\_ MIO chi\_dum | Coef. Std. Err. z P>|z| [95% Conf. Interval] ----age\_range | .0101268 .0042639 2.38 0.018 .0017698 .0184838 hiv5 | -.5955005 .1594579 -3.73 0.000 -.9080323 -.2829687 

Iteration 0: log likelihood = -206.39388
Iteration 1: log likelihood = -202.52284

<sup>.</sup> gen range\_hiv=age\_range\*hiv5

<sup>.</sup> glm chi\_dum age\_range hiv5 range\_hiv, family(binomial n)

log likelihood = -202.522 Iteration 2: Iteration 3: log likelihood = -202.522

Generalized linear models No. of obs 12 Optimization : ML Residual df 8

Scale parameter = 1

Deviance = 345.472226 (1/df) Deviance = 43.18403 (1/df) Pearson = 40.99494= 327.9595101 Pearson

Variance function: V(u) = u\*(1-u/n)[Binomial] Link function : g(u) = ln(u/(n-u))[Logit]

AIC = 34.42033 Log likelihood = -202.5219973BIC = 325.593

chi_dum	   Coef.	OIM Std. Err.	z	P> z		Interval]
age_range hiv5 range_hiv _cons	.0111866	.0043401 .7404394 .0231563 .1277653	2.58 0.48 -1.30 -7.47	0.010 0.631 0.193 0.000	.0026801 -1.095261 075525 -1.20469	.019693 1.807209 .0152462 7038597

. gen age2=(age\_range-27.5)^2

. glm chi\_dum age\_range age2, family(binomial n)

log likelihood = -50.210447Iteration 0: Iteration 1: log likelihood = -50.174929Iteration 2: log likelihood = -50.174929

No. of obs Generalized linear models 12 9 Optimization : ML Residual df Scale parameter = 1

Deviance = 40.77808903 (1/df) Deviance = 4.530899 Pearson = 38.68819948 (1/df) Pearson = 4.298689

Variance function: V(u) = u\*(1-u/n)[Binomial] Link function :  $g(u) = \ln(u/(n-u))$ [Logit]

AIC = 8.862488 Log likelihood = -50.17492879BIC = 18.41393

\_\_\_\_\_\_

	1		OIM						
	chi_dum	Coef.	Std. Err.		P> z	2 - 10	Interval]		
	e_range	.0492848	.0054826	8.99	0.000	.0385391	.0600305		
	age2	0111378	.0006621	-16.82	0.000	0124355	0098401		
	_cons	-1.310733	.1487056	-8.81	0.000	-1.60219	-1.019275		
. glm	. glm chi_dum age_range age2 hiv5, family(binomial n)								
		log likeliho							
1+ara	tion 1.	log likelihe	n = -32.66	31606					

Iteration 0: log likelihood = -32.661696 Iteration 2: log likelihood = -32.661695

 Generalized linear models
 No. of obs = 12

 Optimization : ML
 Residual df = 8

 Scale parameter = 1

 Deviance = 5.751621084
 (1/df) Deviance = .7189526

 Pearson = 6.656666375
 (1/df) Pearson = .8320833

Variance function: V(u) = u\*(1-u/n) [Binomial] Link function : g(u) = ln(u/(n-u)) [Logit]

Log likelihood = -32.66169482 AIC = 6.110282 = -14.12763

| OIM | Chi\_dum | Coef. Std. Err. z P>|z| [95% Conf. Interval] | age\_range | .0538285 .0055535 9.69 0.000 .0429439 .0647132 | age2 | -.0115719 .0006689 -17.30 0.000 -.012883 -.0102608 | hiv5 | -.9203951 .1641225 -5.61 0.000 -1.242069 -.598721 | \_cons | -1.347511 .149278 -9.03 0.000 -1.64009 -1.054931