## How are likelihoods calculated when individual covariates are considered?

Consider a data set for 6 animals:

Y	mass
0	45
1	75
1	80
0	32
1	42
0	50
	Y 0 1 1 0 1 0

A model containing an intercept and mass had the following estimated beta values: b0 = -5.935, b1 = .1146. To calculate the InL for these parameter estimates, you'd use the following equation:

$$\begin{split} \mathbf{L} &= \left[ \frac{e^{(-5.935+0.1146\times45)}}{1+e^{(-5.935+0.1146\times45)}} \right]^0 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times45)}}{1+e^{(-5.935+0.1146\times45)}} \right]^1 \quad \mathbf{x} \\ &\left[ \frac{e^{(-5.935+0.1146\times75)}}{1+e^{(-5.935+0.1146\times75)}} \right]^1 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times75)}}{1+e^{(-5.935+0.1146\times75)}} \right]^0 \quad \mathbf{x} \\ &\left[ \frac{e^{(-5.935+0.1146\times80)}}{1+e^{(-5.935+0.1146\times80)}} \right]^1 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times80)}}{1+e^{(-5.935+0.1146\times80)}} \right]^0 \quad \mathbf{x} \\ &\left[ \frac{e^{(-5.935+0.1146\times32)}}{1+e^{(-5.935+0.1146\times32)}} \right]^0 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times32)}}{1+e^{(-5.935+0.1146\times32)}} \right]^1 \quad \mathbf{x} \\ &\left[ \frac{e^{(-5.935+0.1146\times32)}}{1+e^{(-5.935+0.1146\times42)}} \right]^1 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times32)}}{1+e^{(-5.935+0.1146\times32)}} \right]^1 \quad \mathbf{x} \\ &\left[ \frac{e^{(-5.935+0.1146\times42)}}{1+e^{(-5.935+0.1146\times42)}} \right]^1 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times42)}}{1+e^{(-5.935+0.1146\times42)}} \right]^0 \quad \mathbf{x} \\ &\left[ \frac{e^{(-5.935+0.1146\times42)}}{1+e^{(-5.935+0.1146\times42)}} \right]^1 \times \left[ 1 - \frac{e^{(-5.935+0.1146\times42)}}{1+e^{(-5.935+0.1146\times42)}} \right]^1 \right]^1 \end{split}$$

<u>Remember</u>: if b0 = -5.935, b1 = .1146 are the maximum likelihood estimates, then no other combination of values can be found for b0 and b1 that will have a higher Likelihood value for this dataset and this model.

The calculation on the previous page is simply taking advantage of the following equations, right?

$$\begin{split} pi_{i} &= \frac{e^{\beta_{0} + \beta_{1} \times X1_{i}}}{1 + e^{\beta_{0} + \beta_{1} \times X1_{i}}} \\ L &= \prod_{i=0}^{5} \left( pi_{i} \right)^{y_{i}} \times \left( 1 - pi_{i} \right)^{1 - y_{i}} \\ lL &= \sum_{i=0}^{5} \left[ y_{i} \times \ln(pi_{i}) + \left( 1 - y_{i} \right) \times \ln(1 - pi_{i}) \right] \end{split}$$

For this example, L(b0 = -5.935, b1 = .1146|X) = 0.076The InL = -2.582, and -2InL = 5.165 *Given the preceding equations, do you see why having missing data for only some of the covariates of interest can complicate model comparisons?* 

## Analysis of Mule Deer Fawn Data

Here, you've incorporated information about individual body characteristics. Use of individual covariates greatly expands your ability to model and should be considered for many problems.

Analyses indicated that the best model contains Sex and Length (deltaAIC=0.00)

Ln(S/(1-S) = B0 + B1\*Sex + B2\*Length + B3\*Sex\*Length

LOGIT Link Function Parameters of {11. S(Sex + Length + Length*Sex)										
Parameter	Beta	Standard Error	Lower	Upper						
1: B0 (Intercept)	-22.724138	8.4190916 10.833715	-39.225558							
2: B1 (Sex, Male-1) 3: B3 (length) 4: B4 (sex x length)	0.1858881 -0.1443023	0.0686643	0.0513060	0.3204701						

From looking at the coefficients, we can see that males have a higher intercept (--22.72 + 16.76 = -5.97) and flatter slope (0.186 - 0.144 = 0.04) than do females (intercept = -22.72 and slope = 0.186). So, short females ought to have lower survival than do short males. And, females benefit more than males do from being longer, but ... without calculating actual values for survival and examining them in a table or a figure, it's hard to know much more than that without some pretty hard thinking.

This is where MARK's "Plot Individual Covariates" tool can be very helpful. Choose the model of interest, choose the tool, work with length, and set the sex covariate to either 0 (female) or 1 (male)



So, the point estimates are lower for females at the shorter lengths (but very imprecisely estimated), and, at longer lengths, females appear to be surviving better. A table of values could easily be presented that would show this more concretely.

When we look at such a table, which I created quickly using MARK output, we find that the point estimates are lower for females up to a length of ~116 cm. Beyond that point the females are expected to survive at a higher rate. We could put confidence intervals on the differences using one of several methods designed for such a task. One of these is the delta method, and we'll learn how to use it later in the semester.

	A	В	С	D	E	F	G	Н		J	K	L	М	N	0	P		Q	
1			FEMA	LES			MAL		LES			Difference							
2	Length	1:S	SE	LCI	UCI		1:S	SE	LCI	UCI		Female-Male							
3	108	0.066	0.064	0.009	0.352		0.186	0.144	0.034	0.595		-0.120							
4	108.275	0.069	0.066	0.010	0.355		0.188	0.143	0.036	0.591		-0.118							
5	108.55	0.073	0.068	0.011	0.359		0.189	0.141	0.037	0.587		-0.117							
6	108.825	0.076	0.069	0.012	0.363		0.191	0.140	0.038	0.583		-0.115			I	Female	e-Ⅳ	1ale	
7	109.1	0.080	0.071	0.013	0.366		0.193	0.139	0.040	0.579		-0.113			R				
8	109.375	0.084	0.073	0.014	0.370		0.195	0.138	0.041	0.575		-0.111	0.600						
9	109.65	0.088	0.075	0.015	0.374		0.196	0.136	0.043	0.571		-0.109	0.500					_	
10	109.925	0.092	0.076	0.017	0.377		0.198	0.135	0.045	0.567		-0.106	0.400						
11	110.2	0.096	0.078	0.018	0.381		0.200	0.134	0.046	0.563		-0.104	0.400						
12	110.475	0.101	0.080	0.020	0.385		0.202	0.132	0.048	0.559		-0.101	0.300	-					
13	110.75	0.106	0.081	0.021	0.389		0.204	0.131	0.050	0.555		-0.098	0.200			-			
14	111.025	0.110	0.083	0.023	0.393		0.206	0.130	0.052	0.551		-0.095							
15	111.3	0.116	0.084	0.025	0.397		0.208	0.128	0.054	0.547		-0.092	0.100						
16	111.575	0.121	0.086	0.028	0.400		0.209	0.127	0.056	0.543		-0.088	0.000						
17	111.85	0.126	0.087	0.030	0.404		0.211	0.125	0.058	0.539		-0.085	-0.100	105 110	115	120 1	25	130	135
18	112.125	0.132	0.088	0.033	0.408		0.213	0.124	0.060	0.535		-0.081	0.100						
19	112.4	0.138	0.090	0.035	0.413		0.215	0.122	0.062	0.531		-0.077	-0.200						
20	112.675	0.144	0.091	0.038	0.417		0.217	0.121	0.065	0.527		-0.073							
21	112.95	0.151	0.092	0.042	0.421		0.219	0.119	0.067	0.523		-0.068							
22	113.225	0.157	0.093	0.045	0.425		0.221	0.117	0.069	0.519		-0.063							
23	113.5	0.164	0.094	0.049	0.429		0.223	0.116	0.072	0.515		-0.059							
24	113.775	0.172	0.095	0.053	0.434		0.225	0.114	0.074	0.512		-0.053							
25	114.05	0.179	0.096	0.057	0.438		0.227	0.113	0.077	0.508		-0.048							
26	114.325	0.187	0.096	0.062	0.443		0.229	0.111	0.080	0.504		-0.042							
27	114.6	0.194	0.097	0.067	0.447		0.231	0.109	0.083	0.500		-0.037							
28	114.875	0.203	0.097	0.073	0.452		0.233	0.107	0.086	0.497		-0.030							
29	115.15	0.211	0.097	0.078	0.456		0.235	0.106	0.088	0.493		-0.024							
30	115.425	0.220	0.097	0.085	0.461		0.237	0.104	0.092	0.490		-0.018							
31	115.7	0.228	0.097	0.091	0.466		0.239	0.102	0.095	0.486		-0.011							
32	115.975	0.238	0.097	0.098	0.471		0.241	0.101	0.098	0.483		-0.004							
33	116.25	0.247	0.097	0.106	0.476		0.243	0.099	0.101	0.479		0.004							
34	116.525	0.257	0.096	0.114	0.481		0.246	0.097	0.104	0.476		0.011							
35	116.8	0.267	0.095	0.122	0.486		0.248	0.095	0.108	0.473		0.019							
36	117.075	0.277	0.095	0.131	0.492		0.250	0.093	0.111	0.469		0.027							
37	117.35	0.287	0.094	0.141	0.497		0.252	0.092	0.115	0.466		0.035							
38	117.625	0.298	0.093	0.151	0.503		0.254	0.090	0.119	0.463		0.043							
39	117.9	0.308	0.092	0.161	0.509		0.256	0.088	0.122	0.460		0.052							
40	118.175	0.319	0.090	0.172	0.515		0.258	0.086	0.126	0.457		0.061							
41	118.45	0.331	0.089	0.183	0.521		0.261	0.085	0.130	0.455		0.070							
42	118.725	0.342	0.088	0.195	0.527		0.263	0.083	0.134	0.452		0.079							
43	119	0.354	0.086	0.207	0.534		0.265	0.081	0.138	0.449		0.088							
11	110 275	0.365	0.085	0.220	0.5/11		0.267	11/0	01/11	11/1/	1	n ngg							