

# GENERALIZED LINEAR MODELS FOR BINARY SURVEY VARIABLES

## LOGISTIC REGRESSION

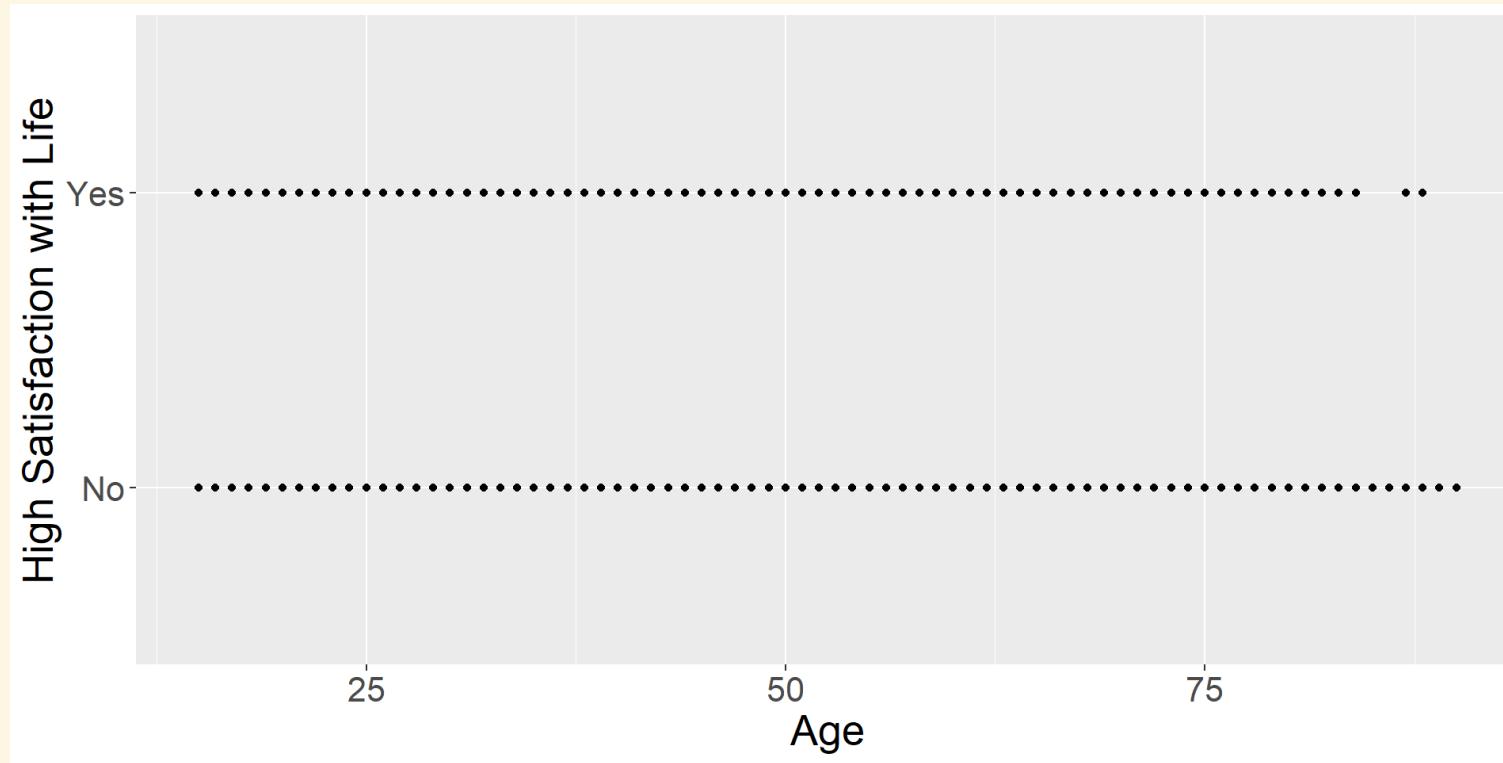
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# WHY LOGISTIC REGRESSION

- When the dependent variable can only be 0 or 1, do we want to draw a straight line through?



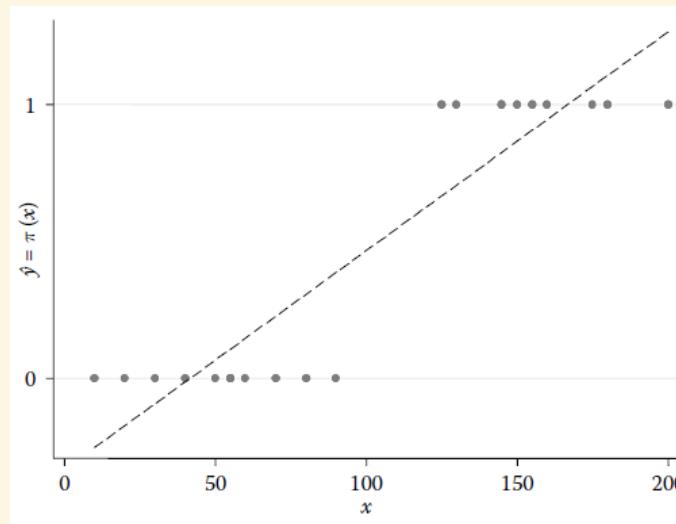
- Rather than predicting these two values, we try to model the *probabilities* that the dependent variable takes one of these two values

# LINEAR TO LOGISTIC REGRESSION

- Consider a linear regression

$$\pi = \Pr(y = 1 | \mathbf{x}) = B_0 + B_1 x_1 + \cdots + B_p x_p + e$$

- The probability must be between 0 and 1, but the linear predictor  $\eta = B_0 + B_1 x_1 + \cdots + B_p x_p$  on the right hand side can take any real number



- Transform the probability to remove the range restrictions, and model the transformation as a linear function of the covariates

# LINEAR TO LOGISTIC REGRESSION

- First, we move from the probability  $[0, 1]$  to the odds  $(0, \infty)$

$$\text{odds} = \frac{\Pr(y = 1|\mathbf{x})}{\Pr(y = 0|\mathbf{x})} = \frac{\pi}{1 - \pi} = e^{B_0 + B_1x_1 + \dots + B_px_p}$$

- Second, we take logarithms to move from the odds to the log-odds  $(-\infty, +\infty)$

$$\text{logit}(\pi) = \ln\left(\frac{\pi}{1 - \pi}\right) = B_0 + B_1x_1 + \dots + B_px_p$$

- Logarithmic transformation maps probabilities from the range  $[0, 1]$  to the entire real line. The probability can be solved from the logit model

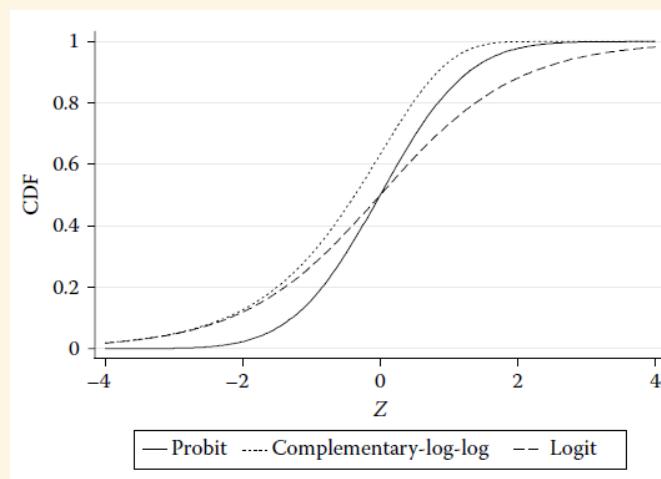
$$\pi = \frac{e^{B_0 + B_1x_1 + \dots + B_px_p}}{1 + e^{B_0 + B_1x_1 + \dots + B_px_p}} = \frac{1}{1 + e^{-(B_0 + B_1x_1 + \dots + B_px_p)}}$$

# LINEAR TO LOGISTIC REGRESSION

- The binary dependent variable  $y$  is assumed to follow a binomial distribution
  - $E(y) = n\pi$
  - $\text{Var}(y) = n\pi(1 - \pi)$
  - The mean and variance depend on the underlying probability  $\pi$
  - Any covariate  $x$  that affects the probability also affects both the mean and variance
- Normality and homoscedasticity assumptions are violated
  - Least squares estimation is not appropriate
  - Maximum likelihood estimation is used

# GENERALIZED LINEAR MODELS

- Generalized linear models have three components
  - Distribution of the dependent variable
  - Linear predictor  $\eta = B_0 + B_1x_1 + \cdots + B_px_p$
  - Link function: the transformation that describes how the mean of the dependent variable is related to the linear predictor  $g(E(y|\mathbf{x})) = \eta$ 
    - Logit  $g(\pi) = \ln(\frac{\pi}{1-\pi})$
    - Probit  $g(\pi) = \Phi^{-1}(\pi)$  where  $\Phi$  is the standard normal cumulative distribution function
    - Complementary log-log  $g(\pi) = \ln(-\ln(1-\pi))$



# MODEL ESTIMATION UNDER SRS

- The logistic regression model is estimated using *maximum likelihood estimation*
- The likelihood function for a SRS of  $n$  independent binomial observations is the product of the probabilities of observing the data given the parameters

$$L(\boldsymbol{\beta}) = \prod_{i=1}^n (\pi_i)^{y_i} (1 - \pi_i)^{1-y_i}$$

- Estimate parameters
  - Take the 1st derivative of  $\ln L(\boldsymbol{\beta})$  with respect to each parameter, set them to zero, and solve for the parameters
  - No closed-form solution, iterative methods (e.g., Newton-Raphson algorithm) are used
- Estimate variance of parameter estimates
  - Take the 2nd derivative of  $\ln L(\boldsymbol{\beta})$  with respect to each parameter, evaluate at the maximum likelihood estimate, and invert to obtain the variance-covariance matrix of the parameter estimates

# MODEL ESTIMATION UNDER COMPLEX SAMPLING

- Maximum likelihood estimation is not appropriate for complex sample designs
  - probability of selection is not constant across observations
  - observations are not independent due to clustering or stratification
- Consider complex sampling from a finite population, *pseudo-maximum likelihood estimation* is used to estimate regression parameters

$$PL(\mathbf{B}) = \prod_{i=1}^n \left( (\pi_i)^{y_i} (1 - \pi_i)^{1-y_i} \right)^{w_i}$$

with

$$\pi_i = \frac{e^{x_i \mathbf{B}}}{1 + e^{x_i \mathbf{B}}}$$

# MODEL ESTIMATION UNDER COMPLEX SAMPLING

- Estimate parameters
  - Maximize the weighted pseudo-likelihood function using the iterative method as in the standard maximum likelihood estimation
- Estimate variance of parameter estimates
  - Taylor series estimation
  - Replication methods (JRR or BRR)

# TESTS OF MODEL PARAMETERS

- Wald test
  - Test the null hypothesis that a parameter is equal to 0
  - The test statistic is the ratio of the parameter estimate to its standard error
  - The test statistic is referred to Student  $t$  distribution with design-based degrees of freedom
  - Alternatively, we can treat the square of the test statistic as a  $\chi^2$  statistic with one degree of freedom
- Likelihood ratio test
  - Compare the likelihood of the model with the parameter of interest to the likelihood of the model without the parameter of interest
  - The test statistic is twice the difference in the log-likelihoods of the two models
  - The test statistic is referred to a  $\chi^2$  distribution with the difference in the number of parameters between the two models
  - *Not applicable* to complex sample designs

# REGRESSION DIAGNOSTICS

- Goodness of Fit (for SRS)
  - Pearson, Deviance
  - Hosmer-Lemeshow test
  - Classification table
  - Area under the ROC curve
  - Psuedo- $R^2$
- Influence and Outliers
  - Cook's distance
  - "Hat" matrix
  - Change in  $\chi^2$  statistic due to deletion of observations

# EXAMPLE: LIFETIME MAJOR DEPRESSION

- Data: National Comorbidity Survey Replication (NCS-R)
- Question: Assess the significance of potential predictors of having lifetime major depression for adults greater than 17 years of age
- Dependent variable:
  - Lifetime major depression (1=Yes; 0=No)
- Predictors:
  - Age (1=18–29; 2=30–44; 3=45–59; 4=60+)
  - Sex (1=Male; 2=Female)
  - Alcohol dependence (1=Yes; 0=No)
  - Education (1=0–11; 2=12; 3=13–15; 4=16+ years)
  - Marital status (1=Married; 2=Previously Married; 3=Never Married)

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## BIVARIATE ANALYSIS

```

1 # Specify survey design
2 ncsrvtyp2 <- svydesign(id = ~seclustr, strata = ~sestrat,
3                           weights = ~ncsrwtlg, data = ncsrp2, nest = TRUE)
4
5 # Bivariate chisq tests
6 svyby(~mdec, ~sexc, design = ncsrvtyp2, svymean)

```

	sexc	mdecNo	mdecYes	se.mdecNo	se.mdecYes
Male	Male	0.8471074	0.1528926	0.009137590	0.009137590
Female	Female	0.7738295	0.2261705	0.006727609	0.006727609

```
1 svychisq(~mdec + sexc, design = ncsrvtyp2)
```

Pearson's  $\chi^2$ : Rao & Scott adjustment

```
data: svychisq(~mdec + sexc, design = ncsrvtyp2)
F = 44.834, ndf = 1, ddf = 42, p-value = 3.965e-08
```

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## ODDS RATIO

	% Having lifetime major depression	Odds
Male	0.1528926	$\frac{0.1528926}{1-0.1528926} = 0.1804879$
Female	0.2261705	$\frac{0.2261705}{1-0.2261705} = 0.2922743$

- Odds ratio

$$\text{OR} = \frac{\text{odds}_{\text{female}}}{\text{odds}_{\text{male}}} = \frac{0.2922743}{0.1804879} = 1.619$$

- The odds of having lifetime major depression are 1.619 times higher for females than for males

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## MODEL SPECIFICATION AND ESTIMATION: SINGLE PREDICTOR

```
1 modell <- svyglm(mdec ~ sexc,
2                     design = ncsrvyp2, family = quasibinomial)
3 summary(modell)
```

Call:

svyglm(formula = mdec ~ sexc, design = ncsrvyp2, family = quasibinomial)

Survey design:

svydesign(id = ~seclustr, strata = ~sestrat, weights = ~ncsrwtlg,  
data = ncsrp2, nest = TRUE)

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-1.71209	0.07055	-24.27	< 2e-16 ***
sexcFemale	0.48203	0.07237	6.66	4.98e-08 ***

---

Signif. codes: 0 '\*\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## INTERPRETATION: SINGLE PREDICTOR

- Logistic regression coefficients represent the change in the log-odds of the dependent variable for a one-unit increase in the predictor, holding all other variables constant
  - The coefficient for *Sex* is 0.48203
  - This means that being female (compared to the reference category, which is male) is associated with an increase in the log-odds of having lifetime major depression
- To make the interpretation more intuitive, we often exponentiate the coefficient to obtain the odds ratio

$$\text{OR} = \frac{\text{odds}_{\text{female}}}{\text{odds}_{\text{male}}} = \frac{e^{B_0+B_1}}{e^{B_0}} = e^{B_1}$$

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## INTERPRETATION: SINGLE PREDICTOR

```
1 exp(model1$coef)
```

```
(Intercept) sexcFemale  
0.1804879 1.6193566
```

- The odds ratio indicates that the odds of having lifetime major depression are 1.619 times higher for females than for males

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## MODEL SPECIFICATION AND ESTIMATION: MULTIPLE PREDICTORS

```

1 model2 <- svyglm(mdec ~ sexc + ag4catc + aldc + ed4catc + mar3catc,
2                     design = ncsrvtyp2, family = quasibinomial)
3 summary(model2)

```

Call:

```
svyglm(formula = mdec ~ sexc + ag4catc + aldc + ed4catc + mar3catc,
       design = ncsrvtyp2, family = quasibinomial)
```

Survey design:

```
svydesign(id = ~seclustr, strata = ~sestrat, weights = ~ncsrwtlg,
          data = ncsrp2, nest = TRUE)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-2.16042	0.15214	-14.200	2.30e-15	***
sexcFemale	0.57735	0.07722	7.477	1.64e-08	***
ag4catc30-44	0.25562	0.09438	2.708	0.0108	*
ag4catc45-59	0.20645	0.09153	2.256	0.0311	*
---	---	---	---	---	---

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## INTERPRETATION: MULTIPLE PREDICTORS

```
1 exp(model2$coef)
```

(Intercept)		sexcFemale	
0.1152765		1.7813032	
ag4catc30-44		ag4catc45-59	
1.2912600		1.2293019	
ag4catc60+		aldcYes	
0.5087563		4.1523575	
ed4catc12		ed4catc13-15	
1.0824803		1.2592434	
ed4catc16+ mar3catcPreviously Married			
1.1769489		1.6264870	
mar3catcNever Married			
1.1225236			

- The odds ratio indicates that the odds of having lifetime major depression are 1.781 times higher for females than for males, holding all other variables constant

# EXAMPLE: LIFETIME MAJOR DEPRESSION

## TESTS OF MODEL PARAMETERS

```
1 # Wald test  
2 regTermTest(model2, ~ag4catc)
```

```
Wald test for ag4catc  
in svyglm(formula = mdec ~ sexc + ag4catc + aldc + ed4catc + mar3catc,  
design = ncsrvtyp2, family = quasibinomial)  
F = 19.98292 on 3 and 32 df: p= 1.7536e-07
```

# SOFTWARE PROCEDURES USING STATA

- The `svy: logit` and `svy: logistic` commands
  - `svy: logistic` defaults to odds ratio output; `coef` option yields logistic model parameter estimates
  - `svy: logit` defaults to log-odds (B) output; or option yields odds ratios “i.” prefix defines categorical predictors
  - Default to lowest alphanumeric category for reference
  - Change reference category for categorical predictors using `ib#.`
  - Post-estimation test statement for Wald tests of multi-parameter hypotheses
    - e.g. (for ASDA Chapter 8 example), to test the null hypothesis that all of the parameters associated with Age are equal to zero, use this test statement:
    - `test 2.ag4cat 3.ag4cat 4.ag4cat`

# SOFTWARE PROCEDURES USING STATA

## BIVARIATE ANALYSIS

```
1 svyset seclustr [pweight = ncsrwtlg], strata(sestrat)
2 svy: tab ag4cat mde, row
3 svy: tab sex mde, row
4 svy: tab ald mde, row
5 svy: tab ed4cat mde, row
6 svy: tab mar3cat mde, row
```

# SOFTWARE PROCEDURES USING STATA

## MODEL SPECIFICATION AND ESTIMATION

```
1 svy: logit mde i.ag4cat ib2.sex ald i.ed4cat i.mar3cat  
2  
3 * Estimated odds ratios and 95% CIs can be generated in svy: logit by addin  
4  
5 svy: logit mde i.ag4cat ib2.sex ald i.ed4cat i.mar3cat, or
```

# SOFTWARE PROCEDURES USING STATA

## WALD TESTS OF MULTI-PARAMETER PREDICTORS

```
1 test 2.ag4cat 3.ag4cat 4.ag4cat  
2 test 2.mar3cat 3.mar3cat  
3 test 2.ed4cat 3.ed4cat 4.ed4cat
```

# SOFTWARE PROCEDURES USING STATA

## TEST OVERALL GOODNESS OF FIT

- Use Archer and Lemeshow's (2006, 2007) design-adjusted test to assess the goodness of fit of this initial model
- `estat gof` (post-estimation command)
- The resulting design-adjusted F-statistic reported in Stata is equal to  $F_{A-L} = 1.229$ , with a p-value of 0.310
  - This suggests that the null hypothesis that the model fits the data well is not rejected
  - We therefore have confidence moving forward that the fit of this initial model is reasonable
- Not presently “canned” in R

# SOFTWARE PROCEDURES USING STATA

## COMPARE LOGIT, PROBIT, AND COMPLEMENTARY LOG-LOG MODELS

```
1 svy: logit ald i.ag4cat ib2.sex i.ed4cat i.mar3cat  
2 svy: probit ald i.ag4cat ib2.sex i.ed4cat i.mar3cat  
3 svy: cloglog ald i.ag4cat ib2.sex i.ed4cat i.mar3cat
```

# SOFTWARE PROCEDURES USING STATA

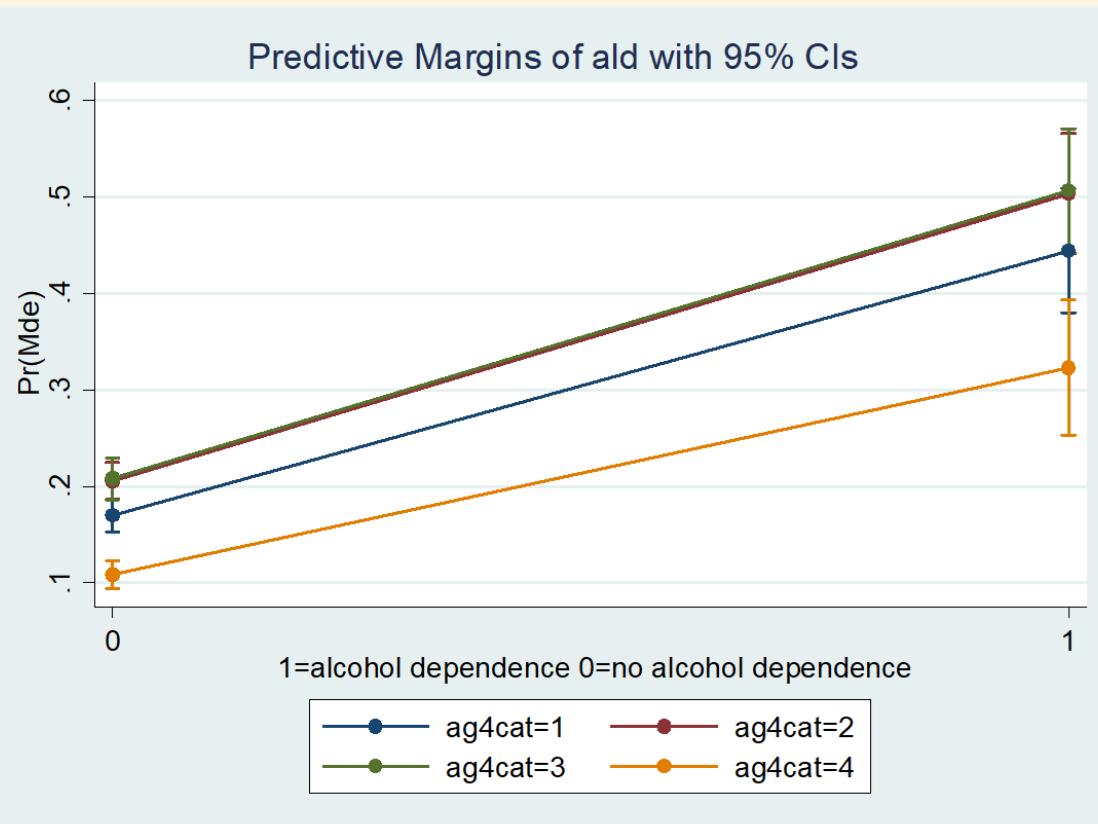
## PLOTTING PREDICTED MARGINAL PROBABILITIES AND EFFECTS

- Stata offers extremely easy-to-use post-estimation commands for calculation and plotting of marginal predicted probabilities based on fitted models (not straightforward in R!)
- Default calculation: compute a model-based predicted probability for everyone in the data set as if they all belonged to the same subgroup, and average the predictions
- One can plot marginal predicted probabilities for different subgroups, or average marginal effects (i.e., expected changes in predicted probabilities associated with a one-unit increase in a given predictor)
- The next few slides present some of the examples illustrated in Chapter 8 of ASDA

# SOFTWARE PROCEDURES USING STATA

## PLOTTING PREDICTED MARGINAL PROBABILITIES AND EFFECTS

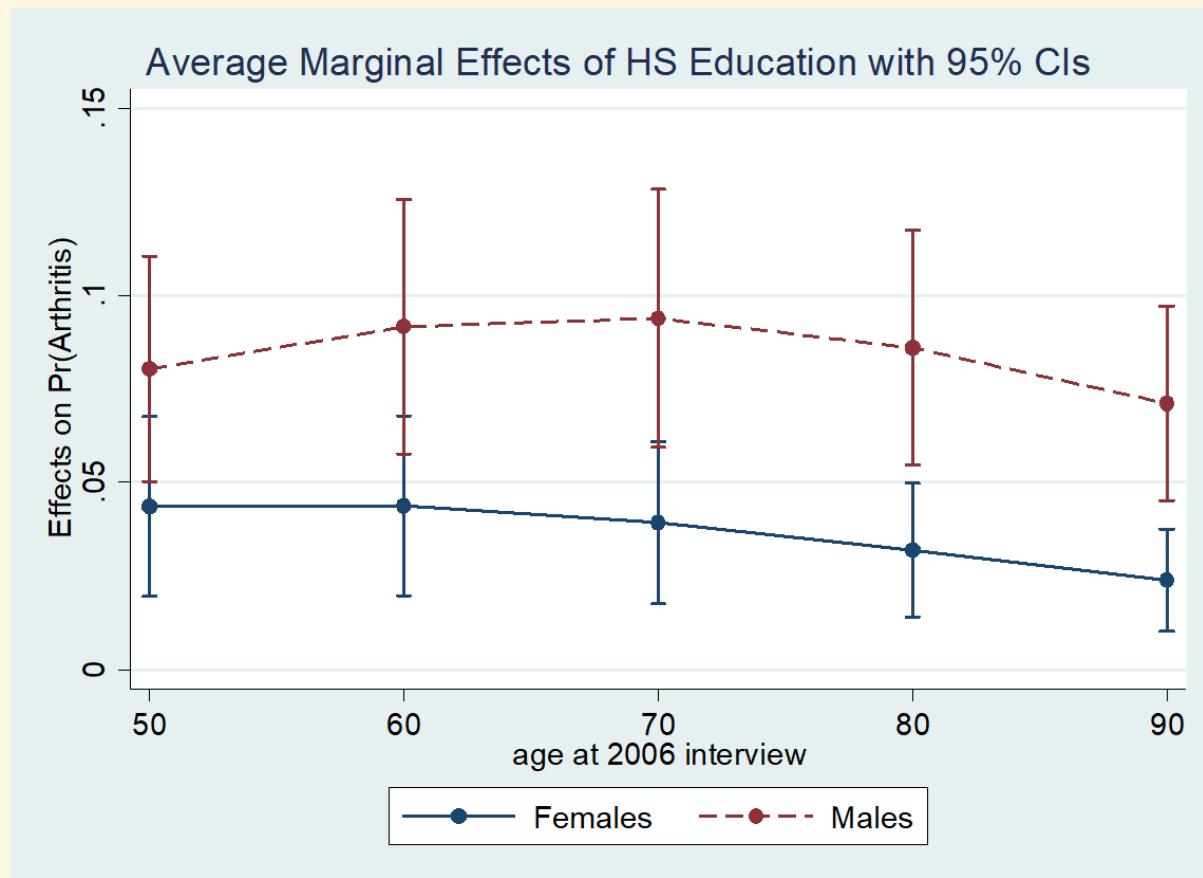
```
1 svy: logit mde i.alld i.ag4cat  
2 margins alld, by(ag4cat)  
3 marginsplot
```



# SOFTWARE PROCEDURES USING STATA

## PLOTTING AVERAGE MARGINAL EFFECTS

```
1 margins, dydx(2.edcat3) by(male) at(kage=(50(10)90))  
2 marginsplot
```



# SOFTWARE PROCEDURES USING STATA

## PLOTTING AVERAGE MARGINAL EFFECTS

```
1 margins, dydx(alld) by(ag4cat)  
2 marginsplot
```

## Average Marginal Effects of ALD with 95% CIs

