

Mixed Models for Longitudinal Binary Outcomes

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Hedeker, D. (2005). Generalized linear mixed models. In B. Everitt & D. Howell (Eds.), *Encyclopedia of Statistics in Behavioral Science*. Wiley.

Hedeker, D. & Gibbons, R.D. (2006). *Longitudinal Data Analysis*, chapter 9. Wiley.

Multilevel models for categorical outcomes

- dichotomous outcomes
 - mixed-effects logistic regression
- ordinal outcomes
 - mixed-effects ordinal logistic regression
 - * proportional odds model
 - * partial or non-proportional odds model
- nominal outcomes
 - mixed-effects nominal logistic regression
- discrete or grouped time-to-event data
 - mixed-effects dichotomous or ordinal regression
 - * complementary log-log link for proportional (& non-proportional) hazards models

www.ssicentral.com/images/pdfs/Survival_clust.pdf

Logistic Regression Model

$$\log \left[\frac{P(Y_i = 1)}{1 - P(Y_i = 1)} \right] = \mathbf{x}'_i \boldsymbol{\beta}$$

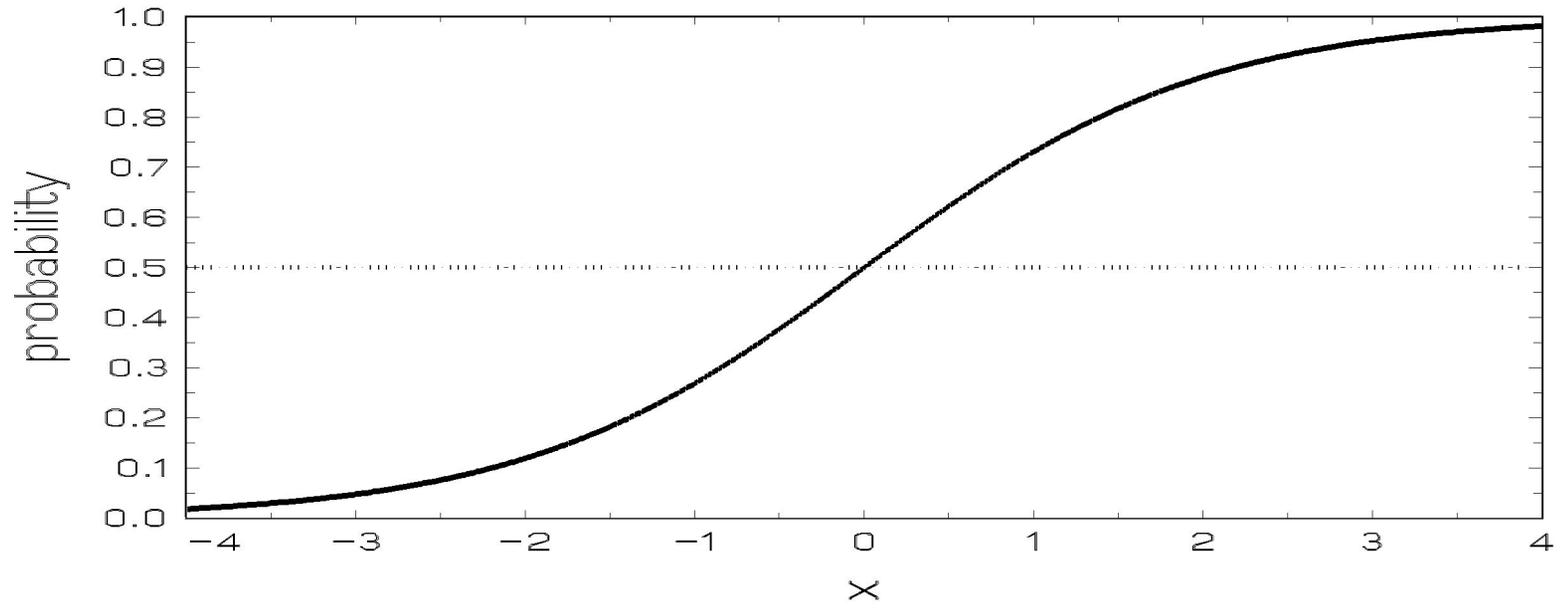
- Dichotomous outcome ($Y = 0$ absence, $Y = 1$ presence).
- Function that links probabilities to regressors is the logit (or log odds) function $\log [P/(1 - P)]$. Logit is called the link function.

The model can be written in terms of probabilities:

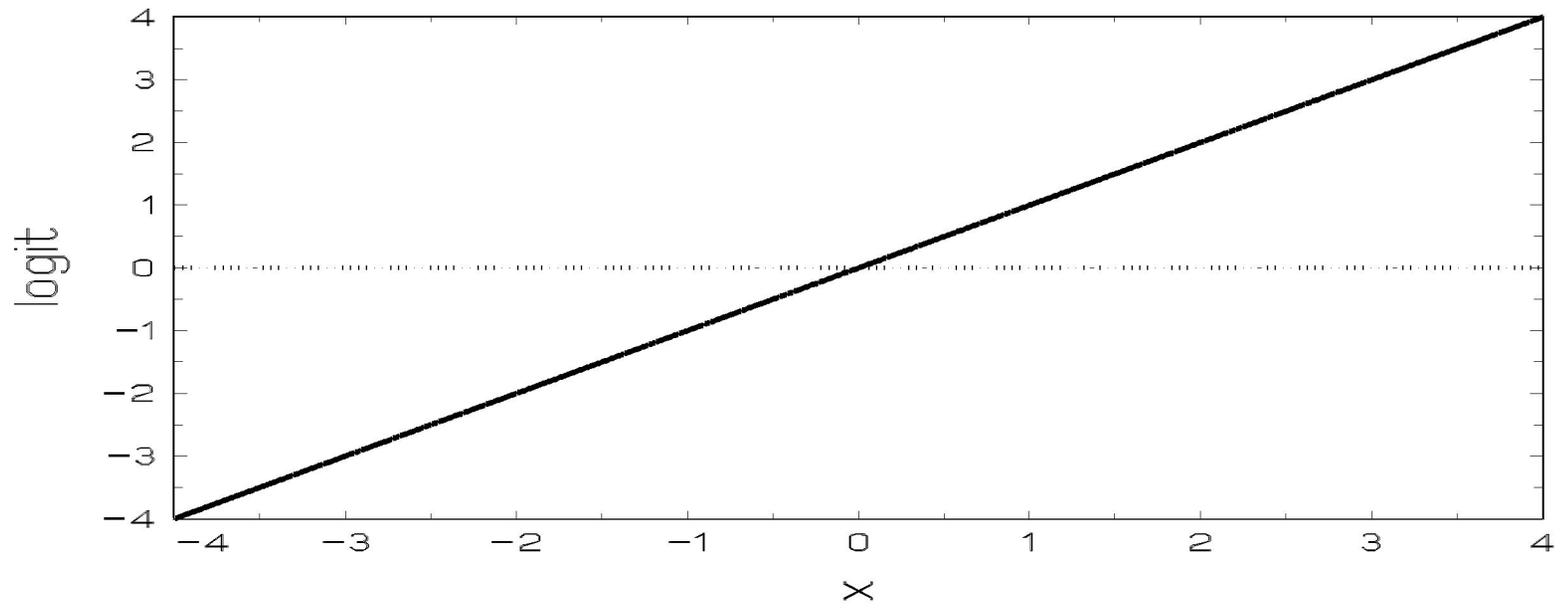
$$P(Y_i = 1) = \frac{1}{1 + \exp(-\mathbf{x}'_i \boldsymbol{\beta})}$$

- Model is a linear model for the logits, not for the probabilities. Logits can take on any values between negative and positive infinity, probabilities can only take on values between 0 and 1.

Logistic Regression Model [slope=1]



Logistic Regression Model [slope=1]



The model can also be written in terms of the odds:

$$\frac{P(Y_i = 1)}{1 - P(Y_i = 1)} = \exp(\mathbf{x}'_i \boldsymbol{\beta})$$

$\exp \beta =$ change in odds for Y per unit change of x

- $\beta = 0$ yields no effect on the odds
- $\beta > 0$ increases odds Y is present with increasing x
- $\beta < 0$ decreases odds Y is present with increasing x

Logistic Regression Model with dichotomous x

group	x	$Y = \text{response}$		prob	odds	logit
		0	1			
control	0	60	30	1/3	1/2	-.693
treatment	1	30	60	2/3	2	.693

$$\log \left[\frac{\text{Pr}(Y_i = 1)}{1 - \text{Pr}(Y_i = 1)} \right] = \beta_0 + \beta_1 x_i$$

$$\exp \beta_0 = \text{odds of response for } x = 0 \quad (30/60 = 1/2)$$

$$\hat{\beta}_0 = \log(1/2) = -.693$$

$$\exp(\beta_0 + \beta_1) = \text{odds of response for } x = 1 \quad (60/30 = 2)$$

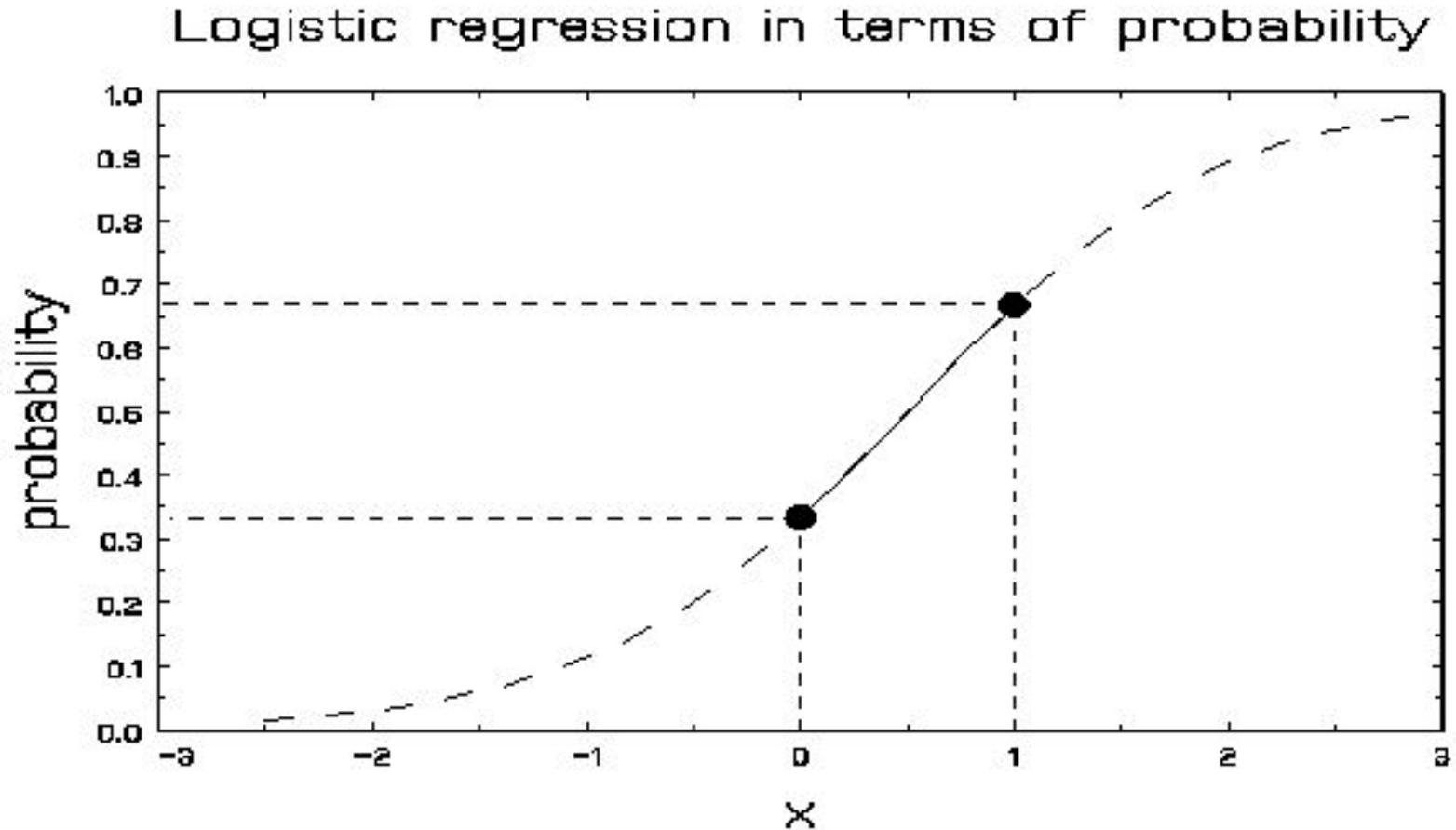
$$\hat{\beta}_0 + \hat{\beta}_1 = \log(2) = .693$$

$$\hat{\beta}_1 = .693 + .693 = 1.386$$

odds ratio = ratio of odds per unit change in x

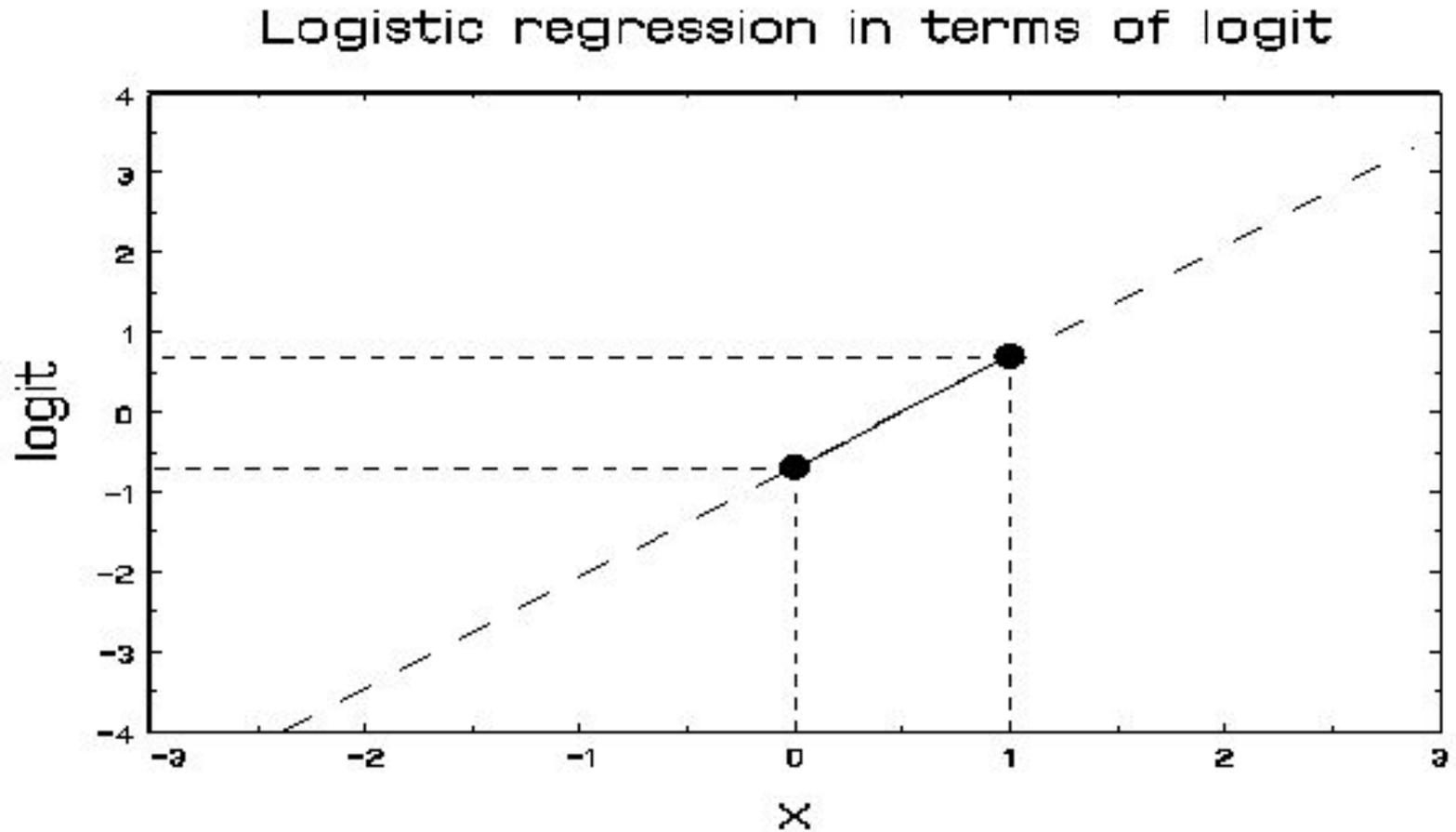
$$\begin{aligned} &= \frac{\exp(\hat{\beta}_0 + \hat{\beta}_1)}{\exp(\hat{\beta}_0)} \\ &= \exp(\hat{\beta}_1) \\ &= \exp(1.386) = 4 \end{aligned}$$

Model is not linear in terms of the probabilities



$$Pr(Y_i = 1) = \frac{1}{1 + \exp[-(\beta_0 + \beta_1 x_i)]} = \frac{\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)}$$

Model is linear in terms of the logits



$$\log \left[\frac{Pr(Y_i = 1)}{1 - Pr(Y_i = 1)} \right] = \beta_0 + \beta_1 x_i$$

Logistic Regression Model with continuous x

age	x	$Y = \text{response}$		prob	odds	logit
		0	1			
20-29	0	60	30	1/3	1/2	-.693
30-39	1	30	60	2/3	2	.693
40-49	2	10	80	8/9	8	2.079

$$\log \left[\frac{\Pr(Y_i = 1)}{1 - \Pr(Y_i = 1)} \right] = \beta_0 + \beta_1 x_i$$

$$\hat{\beta}_0 = -.693$$

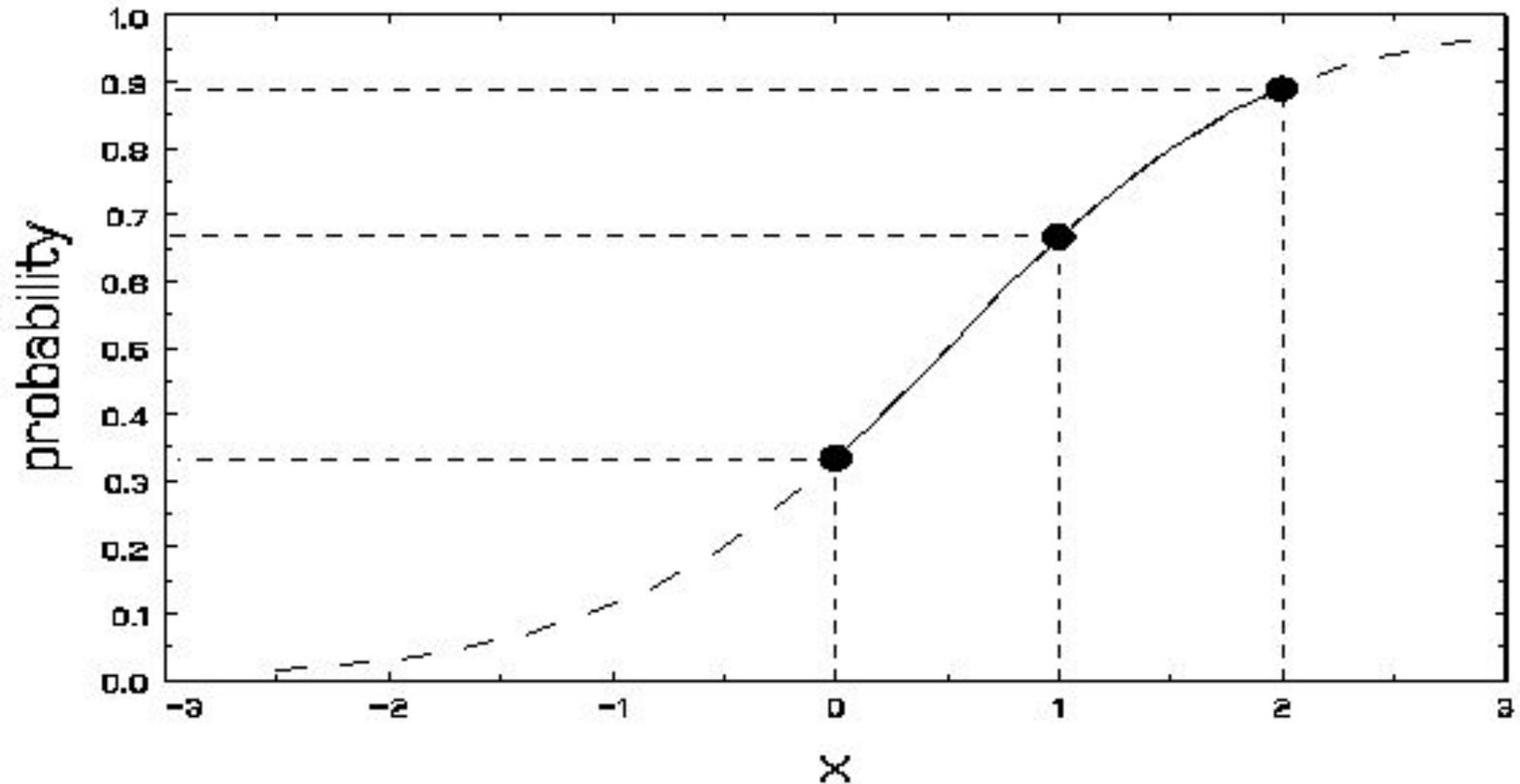
$$\begin{aligned} \hat{\beta}_1 &= \text{change in log odds w/ unit change in } x \\ &= 1.386 \end{aligned}$$

odds ratio = ratio of odds per unit change in x

$$= \exp(\hat{\beta}_1)$$

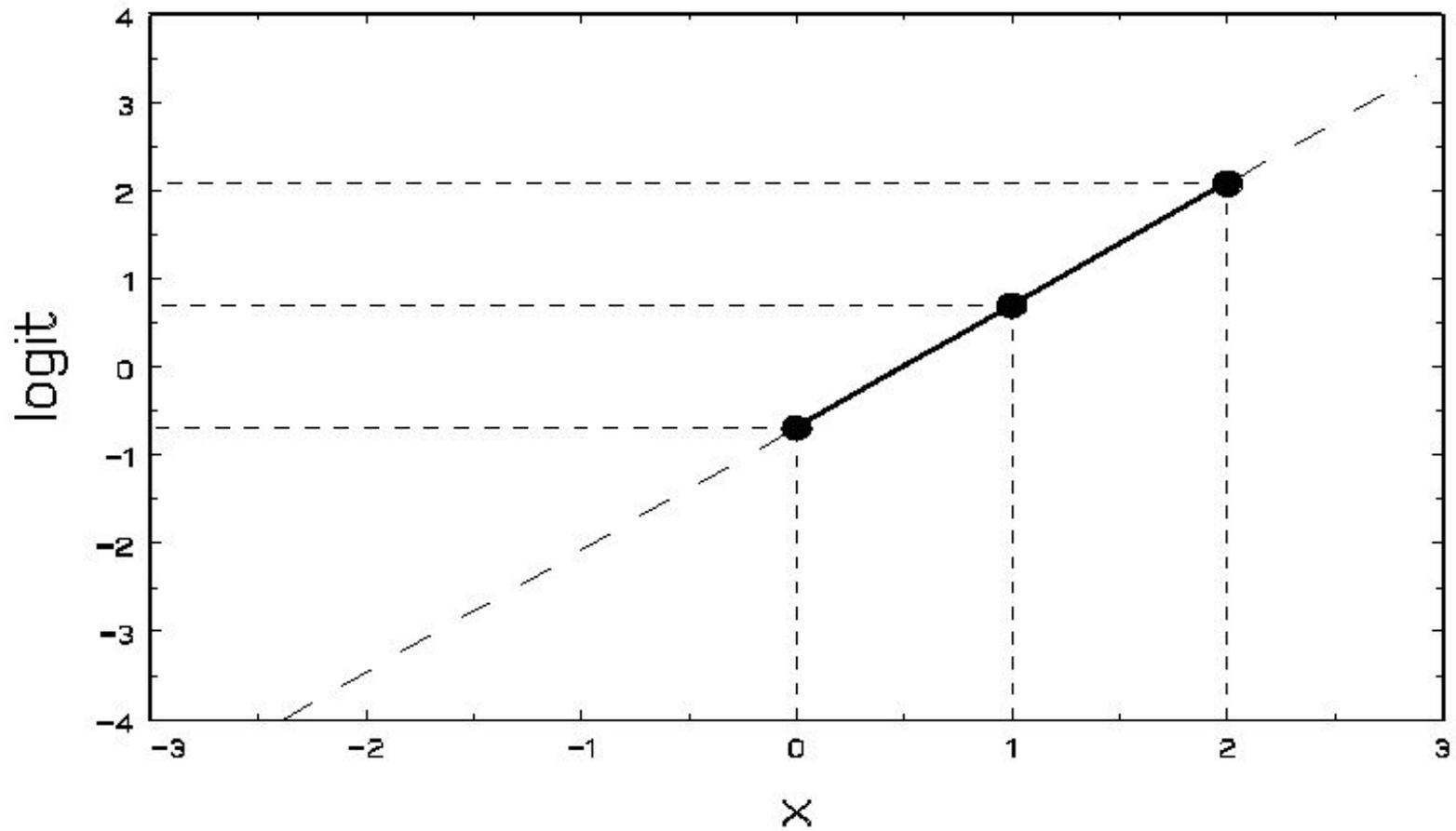
$$= \exp(1.386) = 4$$

Logistic regression in terms of probability



$$Pr(Y_i = 1) = \frac{1}{1 + \exp[-(\beta_0 + \beta_1 x_i)]} = \frac{\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)}$$

Logistic regression in terms of logit

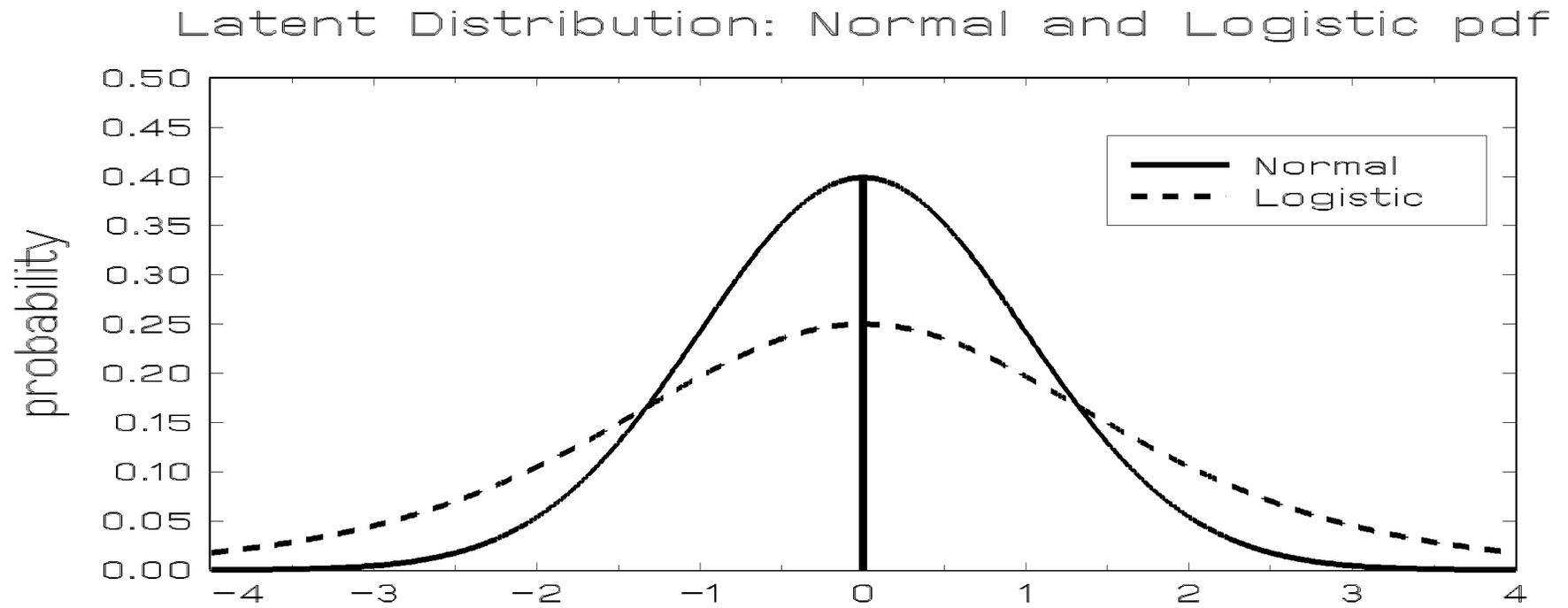


$$\log \left[\frac{Pr(Y_i = 1)}{1 - Pr(Y_i = 1)} \right] = \beta_0 + \beta_1 x_i$$

Dichotomous Response and Threshold Concept

Continuous y_i - an unobservable latent variable - related to dichotomous response Y_i via “threshold concept”

Response occurs ($Y_i = 1$) if $\gamma < y_i$
otherwise, a response does not occur ($Y_i = 0$)



The Threshold Concept in Practice

“How was your day?” (what is your satisfaction level today?)

- Satisfaction may be continuous, but we usually emit a dichotomous response:



Great Day!



a day ...

Model for Latent Continuous Responses

Consider the model with p covariates for the latent response strength y_i ($i = 1, 2, \dots, N$):

$$y_i = \mathbf{x}'_i \boldsymbol{\beta} + \varepsilon_i$$

- probit: $\varepsilon_i \sim$ standard normal (mean=0, variance=1)
 - logistic: $\varepsilon_i \sim$ standard logistic (mean=0, variance= $\pi^2/3$)
- \Rightarrow $\boldsymbol{\beta}$ estimates from logistic regression are larger (in abs. value) than from probit regression by approximately $\sqrt{\pi^2/3} = 1.8$

Underlying latent variable

- useful way of thinking of the problem
- not an essential assumption of the model

Random-intercept Logistic Regression Model

Consider the model with p covariates for the response Y_{ij} for subject i ($i = 1, 2, \dots, N$) at time j ($j = 1, 2, \dots, n_i$):

$$\log \left[\frac{P(Y_{ij} = 1)}{1 - P(Y_{ij} = 1)} \right] = \mathbf{x}'_{ij} \boldsymbol{\beta} + v_{0i}$$

where

Y_{ij} = dichotomous response for subject i at time j

\mathbf{x}_{ij} = $(p + 1) \times 1$ covariate vector (includes 1 for intercept)

$\boldsymbol{\beta}$ = $(p + 1) \times 1$ vector of unknown parameters

v_{0i} = subject effects distributed $\mathcal{NID}(0, \sigma_v^2)$ and assumed independent of \mathbf{x} variables

Model for Latent Continuous Responses

Consider the model with p covariates for the $n_i \times 1$ latent response strength y_{ij} :

$$y_{ij} = \mathbf{x}'_{ij}\boldsymbol{\beta} + \nu_{0i} + \varepsilon_{ij}$$

where assuming

- $\varepsilon_{ij} \sim$ standard normal (mean 0 and $\sigma^2 = 1$) leads to mixed-effects probit regression
- $\varepsilon_{ij} \sim$ standard logistic (mean 0 and $\sigma^2 = \pi^2/3$) leads to mixed-effects logistic regression

Underlying latent variable

- not an essential assumption of the model
- useful for obtaining intra-class correlation (r)

$$r = \frac{\sigma_v^2}{\sigma_v^2 + \sigma^2}$$

and for design effect (d)

$$d = \frac{\sigma_v^2 + \sigma^2}{\sigma^2} = 1/(1 - r)$$

ratio of actual variance to the variance that would be obtained by simple random sampling (holding sample size constant)

Scaling of regression coefficients

β estimates from mixed-effects model are larger (in abs. value) than from fixed-effects model by approximately

$$\sqrt{d} = \sqrt{\frac{\sigma_v^2 + \sigma^2}{\sigma^2}}$$

because

- $V(y) = \sigma_v^2 + \sigma^2$ in mixed-effects model
- $V(y) = \sigma^2$ in fixed-effects model

difference depends on size of random-effects variance σ_v^2

In clustered case, σ_v^2 is usually relatively small and the difference in size of β estimates is also small. However, for longitudinal data, σ_v^2 is usually relatively large and the difference in size of β estimates is therefore much larger.

Treatment-Related Change Across Time

Data from the NIMH Schizophrenia collaborative study on treatment related changes in overall severity. IMPS item 79, *Severity of Illness*, was scored as:

- 1 = normal
- 2 = borderline mentally ill
- 3 = mildly ill

- 4 = moderately ill
- 5 = markedly ill
- 6 = severely ill
- 7 = among the most extremely ill

The experimental design and corresponding sample sizes:

Group	Sample size at Week							<i>completers</i>
	0	1	2	3	4	5	6	
PLC (n=108)	107	105	5	87	2	2	70	65%
DRUG (n=329)	327	321	9	287	9	7	265	81%

Drug = Chlorpromazine, Fluphenazine, or Thioridazine

Main question of interest:

- Was there differential improvement for the drug groups relative to the control group?

Observed proportions \geq “moderately ill”

	<u>week 0</u>	<u>week 1</u>	<u>week 3</u>	<u>week 6</u>
placebo	.98	.91	.89	.71
drug	.99	.82	.66	.42

Observed odds \geq “moderately ill”

	<u>week 0</u>	<u>week 1</u>	<u>week 3</u>	<u>week 6</u>
placebo	52.5	9.50	7.70	2.50
drug	80.8	4.63	1.93	.73
<i>ratio</i>	.65	2.05	3.99	3.42

Observed log odds \geq “moderately ill”

	<u>week 0</u>	<u>week 1</u>	<u>week 3</u>	<u>week 6</u>
placebo	3.96	2.25	2.04	.92
drug	4.39	1.53	.66	-.31
<i>difference</i>	-.43	.72	1.38	1.23
exp (odds ratio)	.65	2.05	3.99	3.42

Stata code: schizb_plots.do

```
log using U:\Data\StatHorizons\schizb_plots.log, replace
infile id imps79 imps79b imps79o inter tx week sweek txswk ///
using https://hedeker.people.uic.edu/SCHIZX1.DAT.txt, clear
recode imps79 imps79b imps79o (-9=.)
```

```
/* only use weeks 0, 1, 3, 6 for these descriptives */
keep if week == 0 | week == 1 | week == 3 | week == 6
summarize
collapse (mean) p1=imps79b, by (tx week)
gen odds1 = p1 / (1-p1)
gen logit1 = log(odds1)
list tx week p1 odds1 logit1
```

```
* graph probabilities & logits
graph twoway (connected p1 week if tx==0) ///
             (connected p1 week if tx==1), ///
             legend(label(1 "placebo") label(2 "drug")) ///
             yscale(range(0 1)) xlabel(0(1)6)
graph twoway (connected logit1 week if tx==0) ///
             (connected logit1 week if tx==1), ///
             legend(label(1 "placebo") label(2 "drug")) ///
             yscale(range(-0.5 4.5)) xlabel(0(1)6)
```

```
gen srweek = sqrt(week)
graph twoway (connected logit1 srweek if tx==0) ///
             (connected logit1 srweek if tx==1), ///
             legend(label(1 "placebo") label(2 "drug")) ///
             yscale(range(-0.5 4.5)) xlabel(0(.5)2.5)
```

```

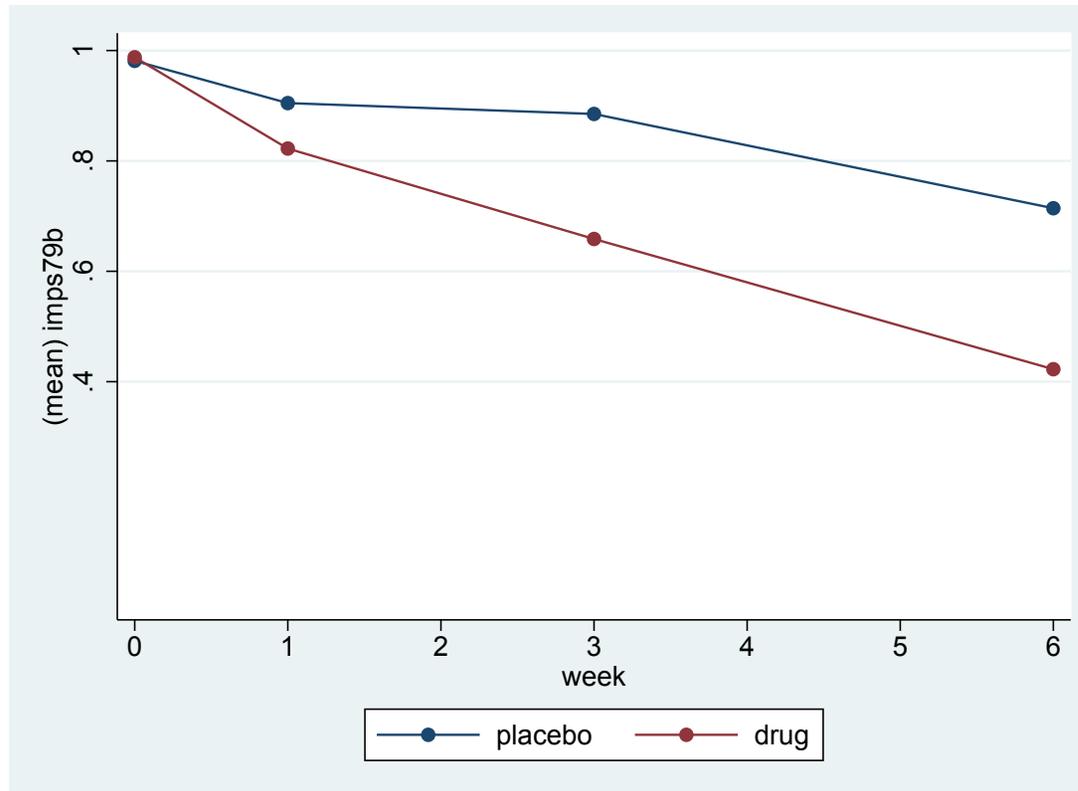
. collapse (mean) p1=imps79b, by (tx week)
. gen odds1 = p1 / (1-p1)
. gen logit1 = log(odds1)
. list tx week p1 odds1 logit1

```

	tx	week	p1	odds1	logit1
1.	0	0	.9813084	52.49997	3.960813
2.	0	1	.9047619	9.500001	2.251292
3.	0	3	.8850574	7.699998	2.04122
4.	0	6	.7142857	2.5	.9162908
5.	1	0	.9877676	80.74995	4.391357
6.	1	1	.8224299	4.631578	1.532898
7.	1	3	.6585366	1.928572	.6567797
8.	1	6	.4226415	.7320262	-.311939

Observed Proportions across Time by Condition

```
. graph twoway (connected p1 week if tx==0) ///  
> (connected p1 week if tx==1), ///  
> legend(label(1 "placebo") label(2 "drug")) ///  
> yscale(range(0 1)) xlabel(0(1)6)
```

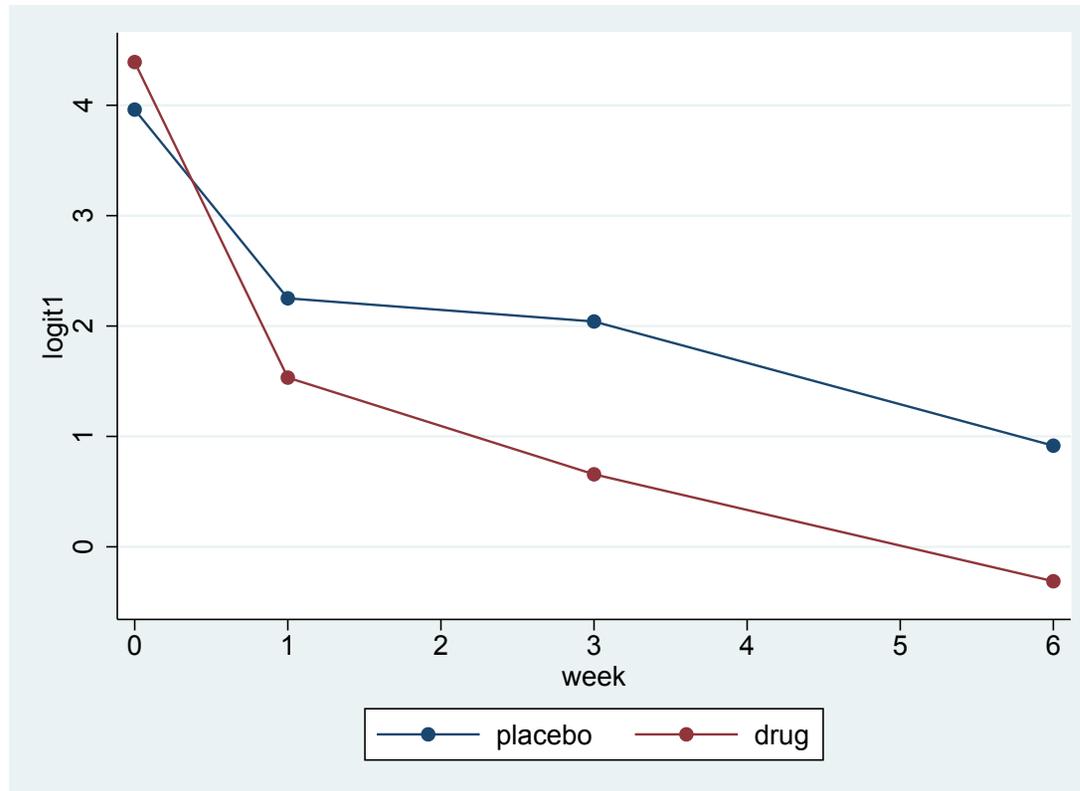


- model is not linear in terms of probabilities

$$Pr(Y_{ij} = 1) = \frac{1}{1 + \exp \left[- \left(\mathbf{x}'_{ij} \boldsymbol{\beta} + v_{0i} \right) \right]}$$

Observed Logits across Time by Condition

```
. graph twoway (connected logit1 week if tx==0) ///  
> (connected logit1 week if tx==1), ///  
> legend(label(1 "placebo") label(2 "drug")) ///  
> yscale(range(-0.5 4.5)) xlabel(0(1)6)
```

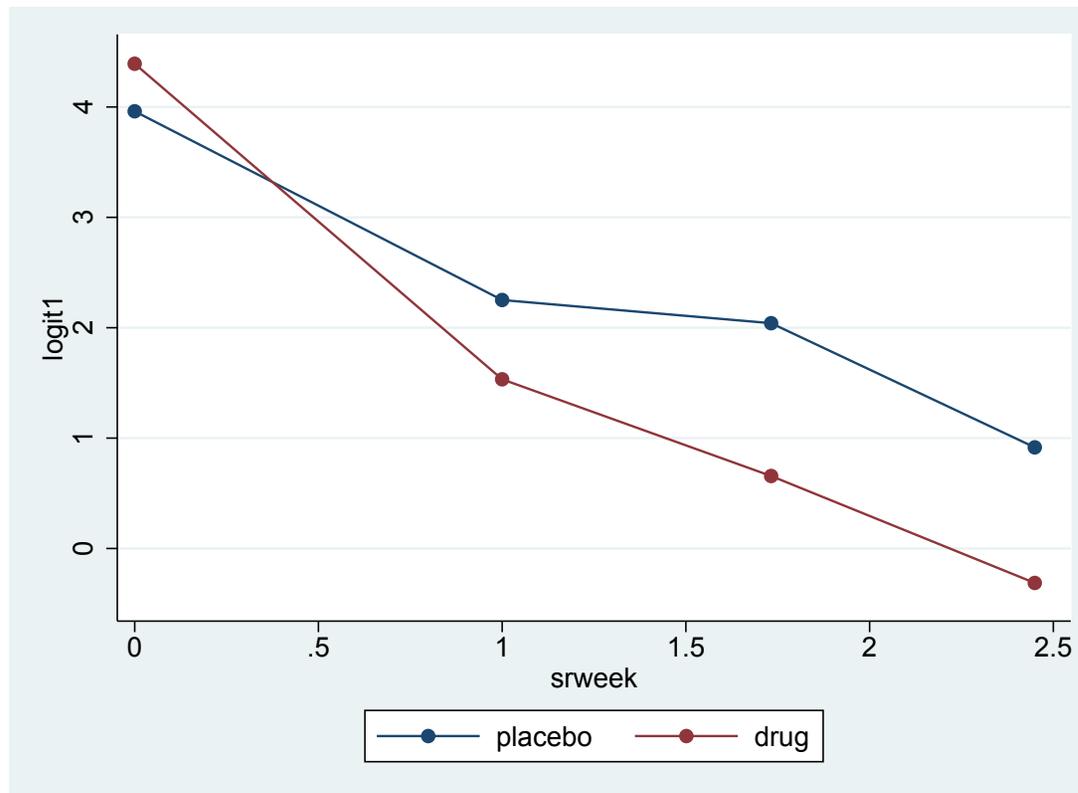


model is linear in terms of logits: $\log \left[\frac{P(Y_{ij} = 1)}{1 - P(Y_{ij} = 1)} \right] = \mathbf{x}'_{ij} \boldsymbol{\beta} + u_{0i}$

```

. gen srweek = sqrt(week)
. graph twoway (connected logit1 srweek if tx==0) ///
> (connected logit1 srweek if tx==1), ///
> legend(label(1 "placebo") label(2 "drug")) ///
> yscale(range(-0.5 4.5)) xlabel(0(.5)2.5)

```



⇒ time effect is approximately linear when expressed as \sqrt{week}

Within-Subjects / Between-Subjects components

Within-subjects model - level 1 ($j = 1, \dots, n_i$ obs)

$$\text{logit}_{ij} = b_{0i} + b_{1i}\sqrt{\text{Week}_j}$$

Between-subjects model - level 2 ($i = 1, \dots, N$ subjects)

$$b_{0i} = \beta_0 + \beta_2 \text{Grp}_i + v_{0i}$$

$$b_{1i} = \beta_1 + \beta_3 \text{Grp}_i$$

$$v_{0i} \sim \mathcal{NID}(0, \sigma_v^2)$$

Put together,

$$\text{logit}_{ij} = (\beta_0 + \beta_2 \text{Grp}_i + v_{0i}) + (\beta_1 + \beta_3 \text{Grp}_i)\sqrt{\text{Week}_j}$$

Stata code: schizb.do

```
log using c:\Stata_Examples\schizb.log, replace
infile id imps79 imps79b imps79o inter tx week sweek txswk ///
using https://hedeker.people.uic.edu/SCHIZX1.DAT.txt, clear
recode imps79 imps79b imps79o (-9=.)
summarize

* random intercept model
melogit imps79b sweek tx txswk || id:
scalar m1 = e(l1)

* random intercept and trend model
melogit imps79b sweek tx txswk || id:sweek, covariance(unstructured)
scalar m2 = e(l1)

* obtain the random effects for each subject
predict u1 u0, reffects

* list the random effects for the first few subjects
by id, sort: generate tolist = (_n==1)
list id u0 u1 in 1/30 if tolist
```

```

* add random and fixed effects together, list out a few
generate intercept = _b[_cons] + _b[tx]*tx + u0
generate slope = _b[sweek] + _b[txswk]*tx + u1
list id intercept slope in 1/30 if tolist

* plot subjects intercepts and slopes
local yref = _b[_cons]
local xref = _b[sweek]
twoway scatter intercept slope, yline('yref', lpattern(dash)) ///
        xline('xref', lpattern(dash))

* generate an estimated spaghetti plot
predict fitimps79b, fitted
sort id sweek
twoway connected fitimps79b sweek, connect(L)

* get LR test for comparing models
display "chibar(12) = " 2*(m2-m1)
display "Prob > chibar2(12) = " .5*chi2tail(1, 2*(m2-m1)) + ///
        .5*chi2tail(2, 2*(m2-m1))
log close

```

```
. summarize
```

Variable	Obs	Mean	Std. Dev.	Min	Max
id	3,059	4777.735	2438.694	1103	9316
imps79	1,603	4.373051	1.471493	1	7
imps79b	1,603	.7679351	.4222819	0	1
imps79o	1,603	2.796007	1.028403	1	4
inter	3,059	1	0	1	1
tx	3,059	.7528604	.4314191	0	1
week	3,059	3	2.000327	0	6
sweek	3,059	1.547414	.7783017	0	2.4495
txswk	3,059	1.164987	.9495875	0	2.4495

Mixed-effects logistic regression
 Group variable: id

Number of obs = 1,603
 Number of groups = 437

Obs per group:
 min = 2
 avg = 3.7
 max = 5

Integration method: mvaghermite
 Integration pts. = 7
 Wald chi2(3) = 170.73
 Log likelihood = -624.82526
 Prob > chi2 = 0.0000

imps79b	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
sweek	-1.498166	.2899992	-5.17	0.000	-2.066554	-.9297781
tx	-.0131162	.6548128	-0.02	0.984	-1.296526	1.270293
txswk	-1.019937	.3345059	-3.05	0.002	-1.675556	-.3643169
_cons	5.383934	.6299046	8.55	0.000	4.149344	6.618524
id						
var(_cons)	4.506713	.9568774			2.972559	6.832652

LR test vs. logistic model: chibar2(01) = 112.41 Prob >= chibar2 = 0.0000

Intraclass Correlation (ICC)

$$ICC = \frac{4.507}{4.507 + \pi^2/3} = 0.578$$

- average pairwise correlation of repeated outcomes = .58
- proportion of (unexplained) variation at the subject level is 58%

Within-Subjects / Between-Subjects components

Within-subjects model - level 1 ($j = 1, \dots, n_i$ obs)

$$\text{logit}_{ij} = b_{0i} + b_{1i}\sqrt{\text{Week}_j}$$

Between-subjects model - level 2 ($i = 1, \dots, N$ subjects)

$$b_{0i} = \beta_0 + \beta_2 \text{Grp}_i + v_{0i}$$

$$b_{1i} = \beta_1 + \beta_3 \text{Grp}_i + v_{1i}$$

$$\mathbf{v}_i \sim \mathcal{NID}(\mathbf{0}, \Sigma_v)$$

Put together,

$$\text{logit}_{ij} = (\beta_0 + \beta_2 \text{Grp}_i + v_{0i}) + (\beta_1 + \beta_3 \text{Grp}_i + v_{1i})\sqrt{\text{Week}_j}$$

```
. melogit imps79b sweek tx txswk || id: sweek, covariance(unstructured)
```

```
Mixed-effects logistic regression      Number of obs      =      1,603  
Group variable:          id           Number of groups   =       437
```

```
Obs per group:  
      min =          2  
      avg =         3.7  
      max =          5
```

```
Integration method: mvaghermite      Integration pts.   =          7
```

```
Wald chi2(3)              =      60.20  
Log likelihood = -613.70418          Prob > chi2       =      0.0000
```

imps79b		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
sweek		-1.429712	.4965533	-2.88	0.004	-2.402939 - .4564858
tx		.2873997	.7558803	0.38	0.704	-1.194098 1.768898
txswk		-1.608433	.4879325	-3.30	0.001	-2.564763 -.6521025
_cons		5.974928	.9836365	6.07	0.000	4.047036 7.902821

id |

var(sweek)	3.093755	1.186109			1.459303	6.558831
var(_cons)	7.226627	3.246852			2.995706	17.433
-----+-----						
id						
cov(_cons,sweek)	-2.172073	1.285726	-1.69	0.091	-4.69205	.3479036

LR test vs. logistic model: chi2(3) = 134.66					Prob > chi2 = 0.0000	

Note: LR test is conservative and provided only for reference.

Postestimation: schizb.do

```
* obtain the random effects for each subject
predict u1 u0, reffects
```

```
* list the random effects for the first few subjects
by id, sort: generate tolist = (_n==1)
list id u0 u1 in 1/30 if tolist
```

```
+-----+
|   id           u0           u1 |
+-----+
1. | 1103   -3.215459   1.264197 |
8. | 1104   -2.318756  -1.059744 |
15. | 1105   -2.349588  -.9809698 |
22. | 1106   -5.687785   .6155943 |
29. | 1107    .4937801   .5770214 |
+-----+
```

```

* add random and fixed effects together, list out a few
generate intercept = _b[_cons] + _b[tx]*tx + u0
generate slope = _b[sweek] + _b[txswk]*tx + u1
list id intercept slope in 1/30 if tolist

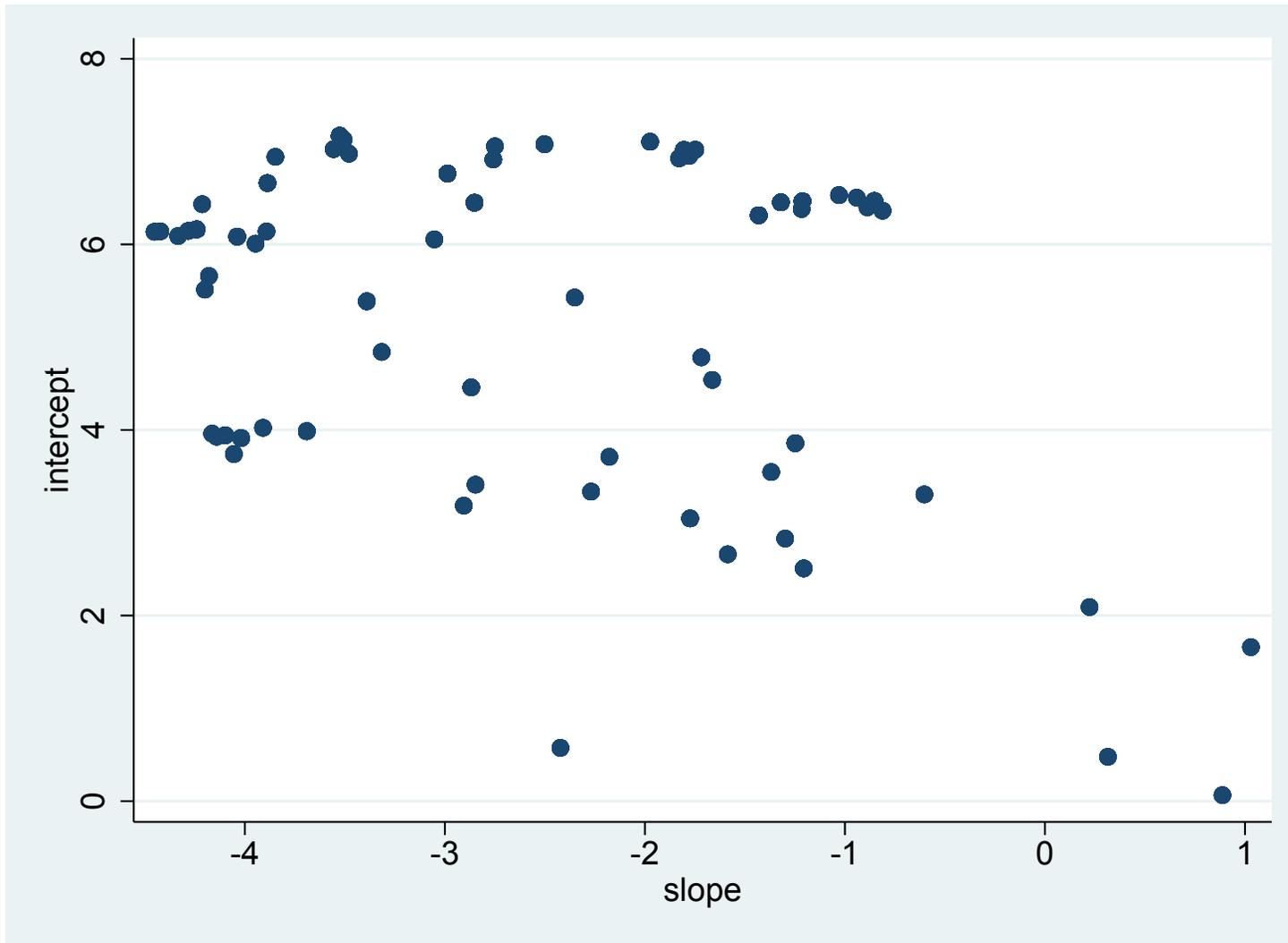
```

```

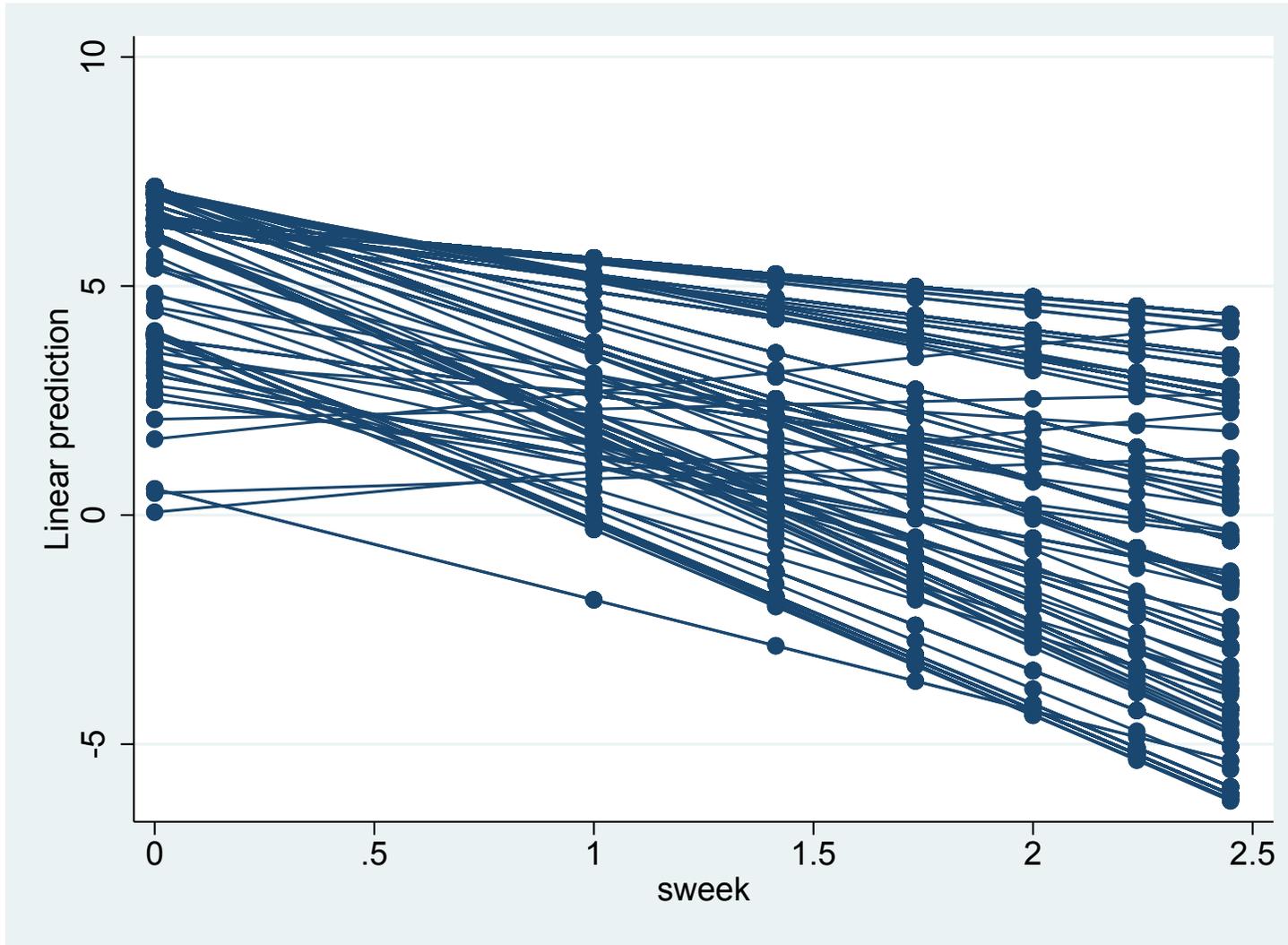
+-----+
|   id   interc~t      slope |
+-----+
1. | 1103   3.046869   -1.773948 |
8. | 1104   3.943572   -4.097889 |
15. | 1105    3.91274   -4.019115 |
22. | 1106    .5745431  -2.422551 |
29. | 1107    6.468709   -.852691 |
+-----+

```

```
* plot subjects intercepts and slopes
local yref = _b[_cons]
local xref = _b[sweek]
twoway scatter intercept slope, yline('yref', lpattern(dash)) ///
      xline('xref', lpattern(dash))
```



```
* generate an estimated spaghetti plot  
predict fitimps79b, fitted  
sort id sweek  
twoway connected fitimps79b sweek, connect(L)
```



Model comparisons - Likelihood Ratio (LR) tests

- comparing random intercept to ordinary logistic regression

LR test vs. logistic model: `chibar2(01) = 112.41 Prob >= chibar2 = 0.0000`

$$H_0 : \sigma_v^2 = 0, \quad H_A : \sigma_v^2 > 0 \quad \Rightarrow \text{one-sided test}$$

`chibar2(01)` refers to a 50:50 mixture of a χ_0^2 and a χ_1^2 distribution; chi-bar square distribution; p -value is obtained from χ_1^2 , but is halved

- comparing random trend to ordinary logistic regression

LR test vs. logistic model: `chi2(3) = 134.66 Prob > chi2 = 0.0000`

Note: LR test is conservative and provided only for reference.

$$H_0 : \sigma_{v_0}^2 = \sigma_{v_1}^2 = \sigma_{v_{01}} = 0$$

- comparing random trend to random intercept model

$$H_0 : \sigma_{v_1}^2 = \sigma_{v_{01}} = 0$$

need `chibar2(12)`, 50:50 mixture of a χ_1^2 and a χ_2^2 distribution; p -value is obtained from the average of χ_1^2 and χ_2^2 (i.e., q and $q - 1$, where q is the number of (co)variance parameters in the null)

```
. * get LR test for comparing models
. display "chibar(12) = " 2*(m2-m1)
chibar(12) = 22.242157

. display "Prob > chibar2(12) = " .5*chi2tail(1, 2*(m2-m1)) + ///
> .5*chi2tail(2, 2*(m2-m1))
Prob > chibar2(12) = 8.600e-06
```

SAS for mixed model analysis of binary outcomes

PROC GLIMMIX (version 9.1.3 and thereafter)

- Multiple levels of nesting, crossed random effects
- Pseudo-likelihood estimation (by default)
 - Linearization to avoid integration over the random effects
 - Produces biased estimates if number of level-1 or level-2 units is small and/or ICC is large
- Full likelihood estimation using numerical quadrature for integration over the random effects **METHOD=QUAD**; however for 3-level models can only use **METHOD=QUAD(QPOINTS=1)** or **METHOD=LAPLACE** (these are equivalent)

PROC NLMIXED

- Full likelihood estimation using numerical quadrature for integration over the random effects
- Only for 2-level models; allows programming features
(can do 3-level models with SAS/STAT 13.2; 2nd maintenance release for SAS 9.4)

SAS Example: schizb.sas

```
FILENAME SchizDat HTTP "https://hedeker.people.uic.edu/SCHIZX1.DAT.txt";  
DATA one; INFILE SchizDat;  
INPUT id imps79 imps79b imps79o int tx week sweek txswk ;
```

sometimes doesn't seem to work

ERROR: The connection has timed out..

NOTE: The SAS System stopped processing this step because of errors.

in this case, easiest just to go to URL and download the data

```
FILENAME SchizDat "U:/mixdemo/SCHIZX1.DAT.txt";  
DATA one; INFILE SchizDat;  
INPUT id imps79 imps79b imps79o int tx week sweek txswk ;
```

SAS code: schizb.sas

```
FILENAME SchizDat HTTP "https://hedeker.people.uic.edu/SCHIZX1.DAT.txt";
DATA one; INFILE SchizDat;
INPUT id imps79 imps79b imps79o int tx week sweek txswk ;

/* get rid of observations with missing values */
IF imps79 > -9;

/* random intercept logistic regression via GLIMMIX */
PROC GLIMMIX DATA=one METHOD=QUAD NOCLPRINT;
CLASS id;
MODEL imps79b(DESC) = tx sweek tx*sweek / SOLUTION DIST=BINARY LINK=LOGIT;
RANDOM INTERCEPT / SUBJECT=id;
COVTEST 'test of random intercept' GLM;
```

- METHOD=QUAD requests full-likelihood estimation (using numerical quadrature)
- COVTEST 'test of random intercept' GLM;
statement yields a likelihood ratio test of
 $H_0 : \sigma_{v_0}^2 = 0, \quad H_A : \sigma_{v_0}^2 > 0 \quad \Rightarrow$ one-sided test

The SAS System

The GLIMMIX Procedure

Model Information

Data Set	WORK.ONE
Response Variable	imps79b
Response Distribution	Binary
Link Function	Logit
Variance Function	Default
Variance Matrix Blocked By	id
Estimation Technique	Maximum Likelihood
Likelihood Approximation	Gauss-Hermite Quadrature
Degrees of Freedom Method	Containment

Number of Observations Read	1603
Number of Observations Used	1603

Response Profile

Ordered Value	imps79b	Total Frequency
1	1	1231
2	0	372

The GLIMMIX procedure is modeling the probability that `imps79b='1'`.

Dimensions

G-side Cov. Parameters	1
Columns in X	4
Columns in Z per Subject	1
Subjects (Blocks in V)	437
Max Obs per Subject	5

Optimization Information

Optimization Technique	Dual Quasi-Newton
Parameters in Optimization	5
Lower Boundaries	1
Upper Boundaries	0
Fixed Effects	Not Profiled
Starting From	GLM estimates
Quadrature Points	7

Iteration History

Iteration	Restarts	Evaluations	Objective Function	Change	Max Gradient
0	0	4	1301.9906335	.	60.97944
1	0	2	1263.9445041	38.04612941	25.65002
2	0	4	1263.3746592	0.56984495	30.57276
.....					

Convergence criterion (GCONV=1E-8) satisfied.

Fit Statistics

-2 Log Likelihood	1249.69
AIC (smaller is better)	1259.69
AICC (smaller is better)	1259.73
BIC (smaller is better)	1280.09
CAIC (smaller is better)	1285.09
HQIC (smaller is better)	1267.74

Fit Statistics for Conditional Distribution

-2 log L(imps79b r. effects)	674.27
Pearson Chi-Square	911.19
Pearson Chi-Square / DF	0.57

Covariance Parameter Estimates

Cov Parm	Subject	Estimate	Standard Error
Intercept	id	4.4825	0.9450

Solutions for Fixed Effects

Effect	Estimate	Standard Error	DF	t Value	Pr > t
Intercept	5.3857	0.6299	435	8.55	<.0001
tx	-0.02140	0.6529	1164	-0.03	0.9739
sweek	-1.4973	0.2905	1164	-5.15	<.0001
tx*sweek	-1.0183	0.3342	1164	-3.05	0.0024

Type III Tests of Fixed Effects

Effect	Num DF	Den DF	F Value	Pr > F
tx	1	1164	0.00	0.9739
sweek	1	1164	26.56	<.0001
tx*sweek	1	1164	9.28	0.0024

Tests of Covariance Parameters
Based on the Likelihood

Label	DF	-2 Log Like	ChiSq	Pr > ChiSq	Note
test of random intercept	1	1362.06	112.37	<.0001	MI

MI: P-value based on a mixture of chi-squares.

Comparing mixed logistic to ordinary (fixed) logistic regression

$$H_0 : \sigma_{v_0}^2 = 0, \quad H_A : \sigma_{v_0}^2 > 0 \quad \Rightarrow \text{one-sided test}$$

mixture refers to a 50:50 mixture of a χ_0^2 and a χ_1^2 distribution;
chi-bar square distribution; p -value is obtained from χ_1^2 , but is halved

```

/* random trend logistic regression via GLIMMIX */
PROC GLIMMIX DATA=one METHOD=QUAD NOCLPRINT;
CLASS id;
MODEL imps79b(DESC) = tx sweek tx*sweek / SOLUTION DIST=BINARY LINK=LOGIT;
RANDOM INTERCEPT sweek / SUBJECT=id TYPE=UN GCORR SOLUTION;
ODS OUTPUT SOLUTIONNR=ebest2;
COVTEST 'test of random effects' GLM;
COVTEST 'test of random slope' . 0 0;

/* print out the estimated random effects */
PROC PRINT DATA=ebest2;
RUN;

```

- **COVTEST 'test of random effects' GLM;**
will compare this model to model without random effects using LR test:
 $H_0 : \sigma_{v_0}^2 = \sigma_{v_1}^2 = \sigma_{v_{01}} = 0$
- **COVTEST 'test of random slope' . 0 0;**
will compare this model to model without random slope and intercept slope covariance using LR test: $H_0 : \sigma_{v_1}^2 = \sigma_{v_{01}} = 0$
- **SOLUTION** on **RANDOM** statement will list the estimates of the random effects

Fit Statistics

-2 Log Likelihood	1227.41
AIC (smaller is better)	1241.41
AICC (smaller is better)	1241.48
BIC (smaller is better)	1269.97
CAIC (smaller is better)	1276.97
HQIC (smaller is better)	1252.68

Estimated G Correlation Matrix

Effect	Row	Col1	Col2
Intercept	1	1.0000	-0.4527
sweek	2	-0.4527	1.0000

Covariance Parameter Estimates

Cov Parm	Subject	Estimate	Standard Error
UN(1,1)	id	6.9816	2.9281
UN(2,1)	id	-2.1027	1.2241
UN(2,2)	id	3.0906	1.1842

Solutions for Fixed Effects

Effect	Estimate	Standard Error	DF	t Value	Pr > t
Intercept	5.9323	0.9481	435	6.26	<.0001
tx	0.2882	0.7436	727	0.39	0.6985
sweek	-1.4042	0.4734	437	-2.97	0.0032
tx*sweek	-1.6115	0.4806	727	-3.35	0.0008

Solution for Random Effects

Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t
Intercept	id 1103	-3.5747	2.1974	727	-1.63	0.1042
sweek	id 1103	1.5078	1.2532	727	1.20	0.2293
Intercept	id 1104	-2.4664	2.1680	727	-1.14	0.2556
sweek	id 1104	-0.7525	1.6193	727	-0.46	0.6423
Intercept	id 1105	-2.4732	2.1699	727	-1.14	0.2548
sweek	id 1105	-0.7363	1.6370	727	-0.45	0.6530

Tests of Covariance Parameters
Based on the Likelihood

Label	DF	-2 Log Like	ChiSq	Pr > ChiSq	Note
test of random effects	3	1362.06	134.66	<.0001	--
test of random slope	2	1249.73	22.33	<.0001	MI

MI: P-value based on a mixture of chi-squares.

--: Standard test with unadjusted p-values.

- **test of random effects** compares random trend model to ordinary logistic regression using ordinary LR test (too conservative)
- **test of random slope** compares random trend to random intercept model

$$H_0 : \sigma_{v_1}^2 = \sigma_{v_{01}} = 0$$

uses `chibar2(12)`, 50:50 mixture of a χ_1^2 and a χ_2^2 distribution; p -value is obtained from the average of χ_1^2 and χ_2^2 (i.e., q and $q - 1$, where q is the number of (co)variance parameters in the null)

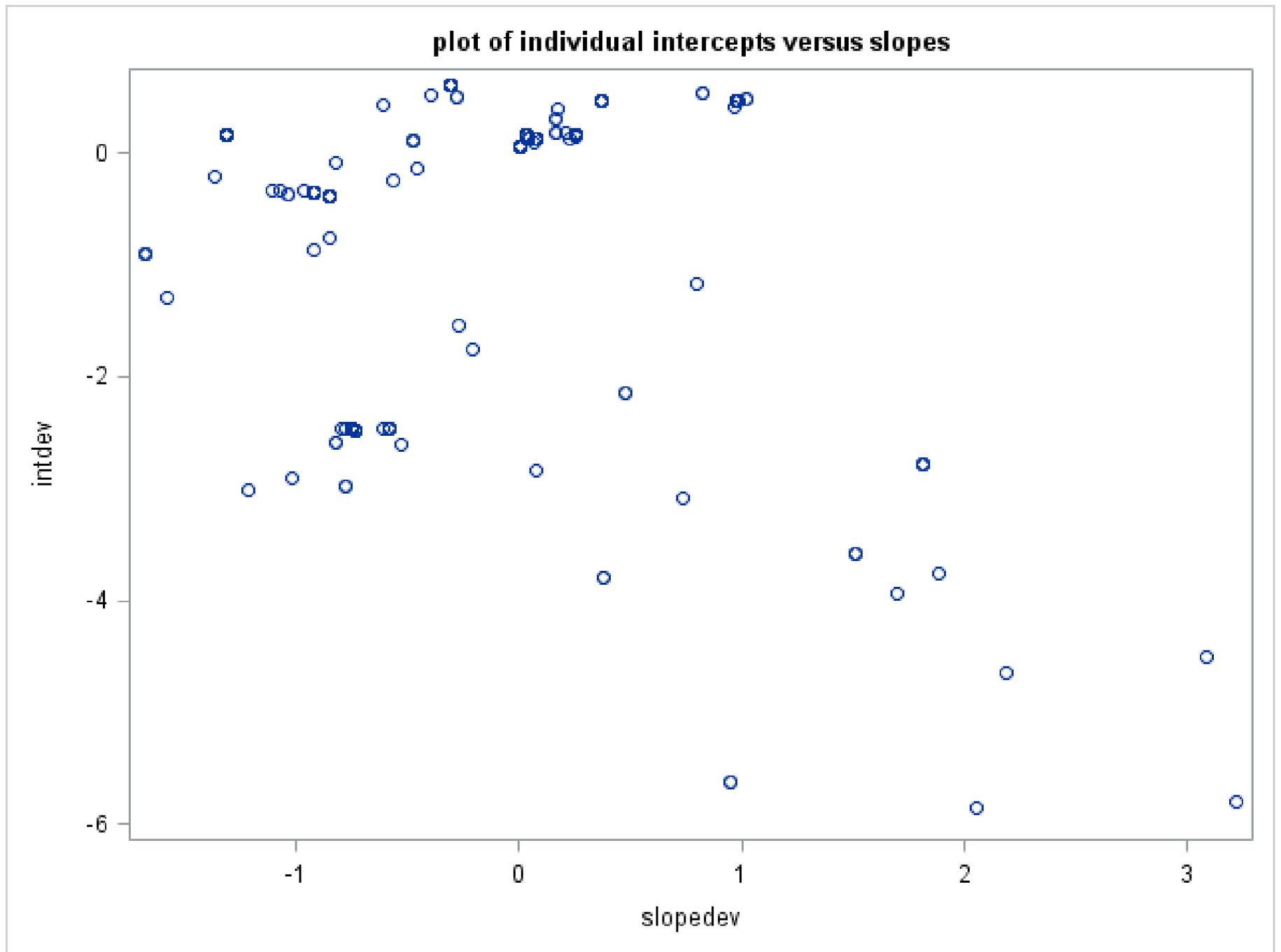
```

/* print out the estimated random effects */
PROC PRINT DATA=ebest2;
RUN;

```

Obs	Effect	Subject	Estimate	StdErr Pred	DF	tValue	Probt
1	Intercept	id 1103	-3.5747	2.1974	727	-1.63	0.1042
2	sweek	id 1103	1.5078	1.2532	727	1.20	0.2293
3	Intercept	id 1104	-2.4664	2.1680	727	-1.14	0.2556
4	sweek	id 1104	-0.7525	1.6193	727	-0.46	0.6423
5	Intercept	id 1105	-2.4732	2.1699	727	-1.14	0.2548
6	sweek	id 1105	-0.7363	1.6370	727	-0.45	0.6530
7	Intercept	id 1106	-5.6168	1.9482	727	-2.88	0.0041
8	sweek	id 1106	0.9518	1.4954	727	0.64	0.5247

```
DATA randest2 (KEEP=subject intdev slopedev);  
  ARRAY y(2) intdev slopedev;  
  DO par = 1 TO 2;  
    SET ebest2;  
    BY subject;  
    y(par) = ESTIMATE;  
    IF LAST.id THEN RETURN;  
  END;  
  
PROC SGPLOT DATA=randest2;  
  SCATTER X=slopedev Y=intdev;  
  TITLE2 'plot of individual intercepts versus slopes';  
RUN;
```



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Instructions

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- Open SuperMixTrialLicense.xls. Enter the personal data requested in the first row of SuperMixTrialLicense.xls into the corresponding columns of the second row of SuperMixTrialLicense.xls. Save the revised copy of SuperMixTrialLicense.xls.
- Compose an email message to SSI Customer Support with the subject line SuperMix Trial License.
- Attach the saved copy of SuperMixTrialLicense.xls to the email message to SSI Customer Support.
- Send the email message to [SSI Customer Support](#).

Free student edition

A student version of SuperMix for Windows can be downloaded from this page.

Please print this page before downloading the student edition, as the restrictions and other useful information concerning the program are given here. Also note that **no technical support is provided** for the student edition and that no serial number is required during installation.

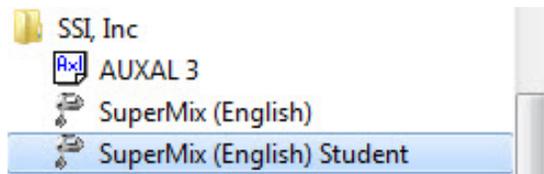
Installation:

SUPERMIX - MENU

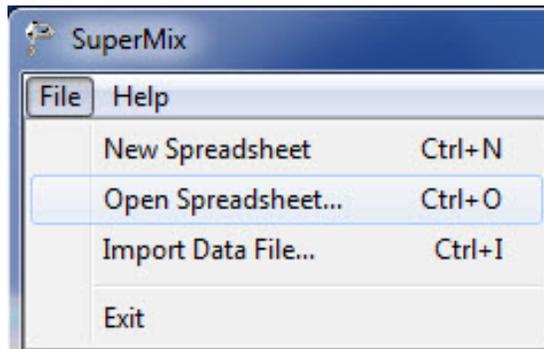
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www.ssicentral.com/index.php/products/supermix/downloads

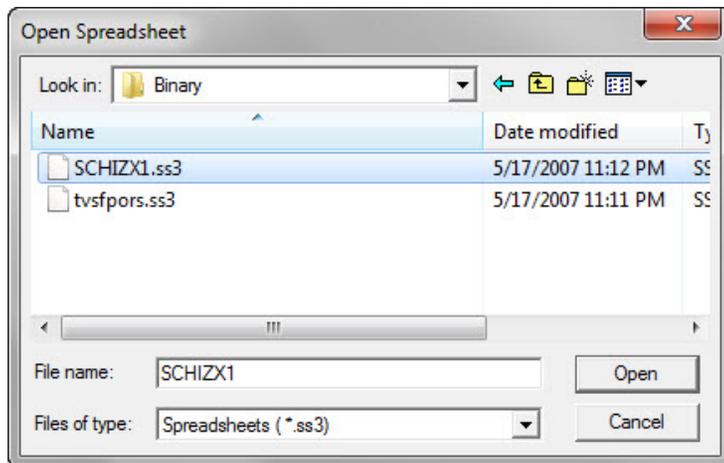
- Under SSI, Inc > “SuperMix (English)” or “SuperMix (English) Student”



- Under “File” click on “Open Spreadsheet”



- Open C:\SuperMixEn Examples\Workshop\Binary\SCHIZX1.ss3
(or C:\SuperMixEn Student Examples\Workshop\Binary\SCHIZX1.ss3)



C:\SuperMixEn Examples\Workshop\Binary\SCHIZX1.ss3

SCHIZX1.ss3

1103 Apply

	(A)_Patient	(B)_Imps79	(C)_Imps79D	(D)_Imps79D	(E)_TxDrug	(F)_Week	(G)_SqrtWee	(H)_Tx*SWe
1	1103	5.50	1	4	1	0	0.00	0.00
2	1103	3.00	0	2	1	1	1.00	1.00
3	1103	-9.00	-9	-9	1	2	1.41	1.41
4	1103	2.50	0	2	1	3	1.73	1.73
5	1103	-9.00	-9	-9	1	4	2.00	2.00
6	1103	-9.00	-9	-9	1	5	2.24	2.24
7	1103	4.00	1	2	1	6	2.45	2.45
8	1104	6.00	1	4	1	0	0.00	0.00
9	1104	3.00	0	2	1	1	1.00	1.00
10	1104	-9.00	-9	-9	1	2	1.41	1.41
11	1104	1.50	0	1	1	3	1.73	1.73
12	1104	-9.00	-9	-9	1	4	2.00	2.00
13	1104	-9.00	-9	-9	1	5	2.24	2.24
14	1104	2.50	0	2	1	6	2.45	2.45
15	1105	4.00	1	2	1	0	0.00	0.00
16	1105	3.00	0	2	1	1	1.00	1.00
17	1105	-9.00	-9	-9	1	2	1.41	1.41
18	1105	1.00	0	1	1	3	1.73	1.73
19	1105	-9.00	-9	-9	1	4	2.00	2.00
20	1105	-9.00	-9	-9	1	5	2.24	2.24
21	1105	-9.00	-9	-9	1	6	2.45	2.45
22	1106	3.00	0	2	1	0	0.00	0.00
23	1106	1.00	0	1	1	1	1.00	1.00
24	1106	-9.00	-9	-9	1	2	1.41	1.41
25	1106	1.50	0	1	1	3	1.73	1.73
26	1106	-9.00	-9	-9	1	4	2.00	2.00
27	1106	-9.00	-9	-9	1	5	2.24	2.24
28	1106	1.00	0	1	1	6	2.45	2.45

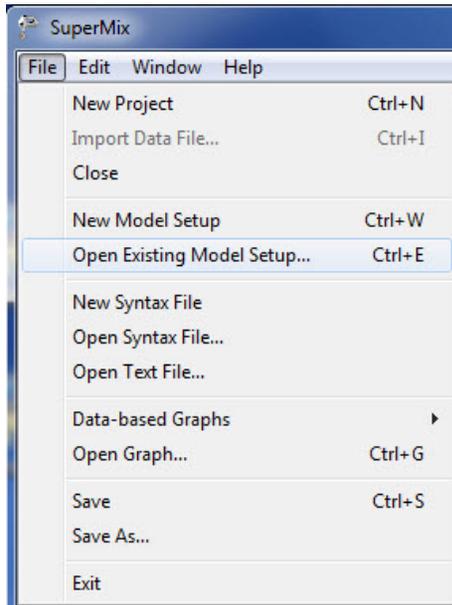
Select Imps79D column, then “Edit” > “Set Missing Value”

The screenshot shows a data editor window titled "SCHIZX1.ss3" with a table of 28 rows and 8 columns. The columns are labeled (A)_Patient, (B)_Imps79, (C)_Imps79D, (D)_Imps79D, (E)_TxDrug, (F)_Week, (G)_SqrtWee, and (H)_Tx*SWe. The data in the table is as follows:

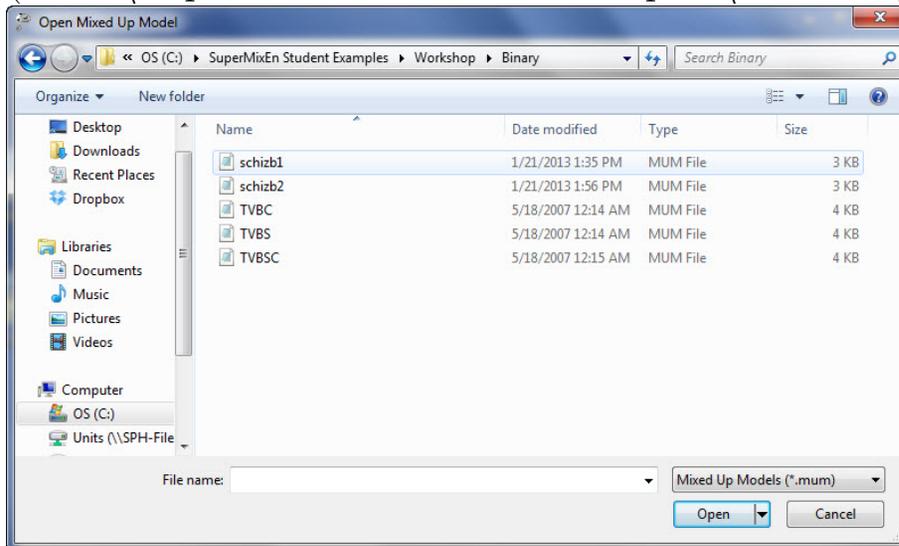
	(A)_Patient	(B)_Imps79	(C)_Imps79D	(D)_Imps79D	(E)_TxDrug	(F)_Week	(G)_SqrtWee	(H)_Tx*SWe
1	1103	5.50	1	4	1	0	0.00	0.00
2	1103	3.00	0	2	1	1	1.00	1.00
3	1103	-9.00	-9	-9	1	2	1.41	1.41
4	1103	2.50	0	2	1	3	1.73	1.73
5	1103	-9.00	-9	-9	1	4	2.00	2.00
6	1103	-9.00	-9	-9	1	5	2.24	2.24
7	1103	4.00	1	2	1	6	2.45	2.45
8	1104	6.00	1	4	1	0	0.00	0.00
9	1104	3.00	0	2	1	1	1.00	1.00
10	1104	-9.00	-9	-9	1	2	1.41	1.41
11	1104	1.50	0	1	1	3	1.73	1.73
12	1104	-9.00	-9	-9	1	4	2.00	2.00
13	1104	-9.00	-9	-9	1	5	2.24	2.24
14	1104	2.50	0	2	1	6	2.45	2.45
15	1105	4.00	1	2	1	0	0.00	0.00
16	1105	3.00	0	2	1	1	1.00	1.00
17	1105	-9.00	-9	-9	1	2	1.41	1.41
18	1105	1.00	0	1	1	3	1.73	1.73
19	1105	-9.00	-9	-9	1	4	2.00	2.00
20	1105	-9.00	-9	-9	1	5	2.24	2.24
21	1105	-9.00	-9	-9	1	6	2.45	2.45
22	1106	3.00	0	2	1	0	0.00	0.00
23	1106	1.00	0	1	1	1	1.00	1.00
24	1106	-9.00	-9	-9	1	2	1.41	1.41
25	1106	1.50	0	1	1	3	1.73	1.73
26	1106	-9.00	-9	-9	1	4	2.00	2.00
27	1106	-9.00	-9	-9	1	5	2.24	2.24
28	1106	1.00	0	1	1	6	2.45	2.45

A dialog box is open over the table, showing "Missing Value Code: -9" with "OK" and "Cancel" buttons.

Under “File” click on “Open Existing Model Setup”



Open C:\SuperMixEn Examples\Workshop\Binary\schizb1.mum
(or C:\SuperMixEn Student Examples\Workshop\Binary\schizb1.mum)



Note “Dependent Variable Type” should be “binary”

Model Setup: schizb1.mum

Configuration | Variables | Starting Values | Patterns | Advanced | Linear Transforms

Title 1: Schiz BINARY outcome

Title 2: random intercept model

Dependent Variable Type: **binary**

Level-2 ID: Patient

Dependent Variable: Imps79D

Level-3 ID:

Write Bayes Estimates: no

Convergence Criterion: 0.0001

Number of Iterations: 100

Categories:

	Value
1	0
2	1

Missing Values Present: true

Perform Crosstabulation: no

Missing Value for the Dependent Var: -9.0

Global Missing Value: -9.0

Output Type: standard

Select the form of the dependent variable. The options on the screens will change as required.

Model Setup: schizb1.mum

Configuration | **Variables** | Starting Values | Patterns | Advanced | Linear Transforms

Available	E	2
Patient	<input type="checkbox"/>	<input type="checkbox"/>
Imps79	<input type="checkbox"/>	<input type="checkbox"/>
Imps79D	<input type="checkbox"/>	<input type="checkbox"/>
Imps790	<input type="checkbox"/>	<input type="checkbox"/>
TxDrug	<input checked="" type="checkbox"/>	<input type="checkbox"/>
Week	<input type="checkbox"/>	<input type="checkbox"/>
SqrtWeek	<input checked="" type="checkbox"/>	<input type="checkbox"/>
Tx*SWeek	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Explanatory Variables

- TxDrug
- SqrtWeek
- Tx*SWeek

L-2 Random Effects

Include Intercept

Include Intercept

Use the arrow keys or click on the desired tab to select the category of interest for the model.

Note “Optimization Method” should be “adaptive quadrature”

Model Setup: schizb1.mum

Configuration | Variables | Starting Values | Patterns | **Advanced** | Linear Transforms

General Settings

Unit Weighting: equal

Optimization Method: **adaptive quadrature**

Number of Quadrature Points: 25

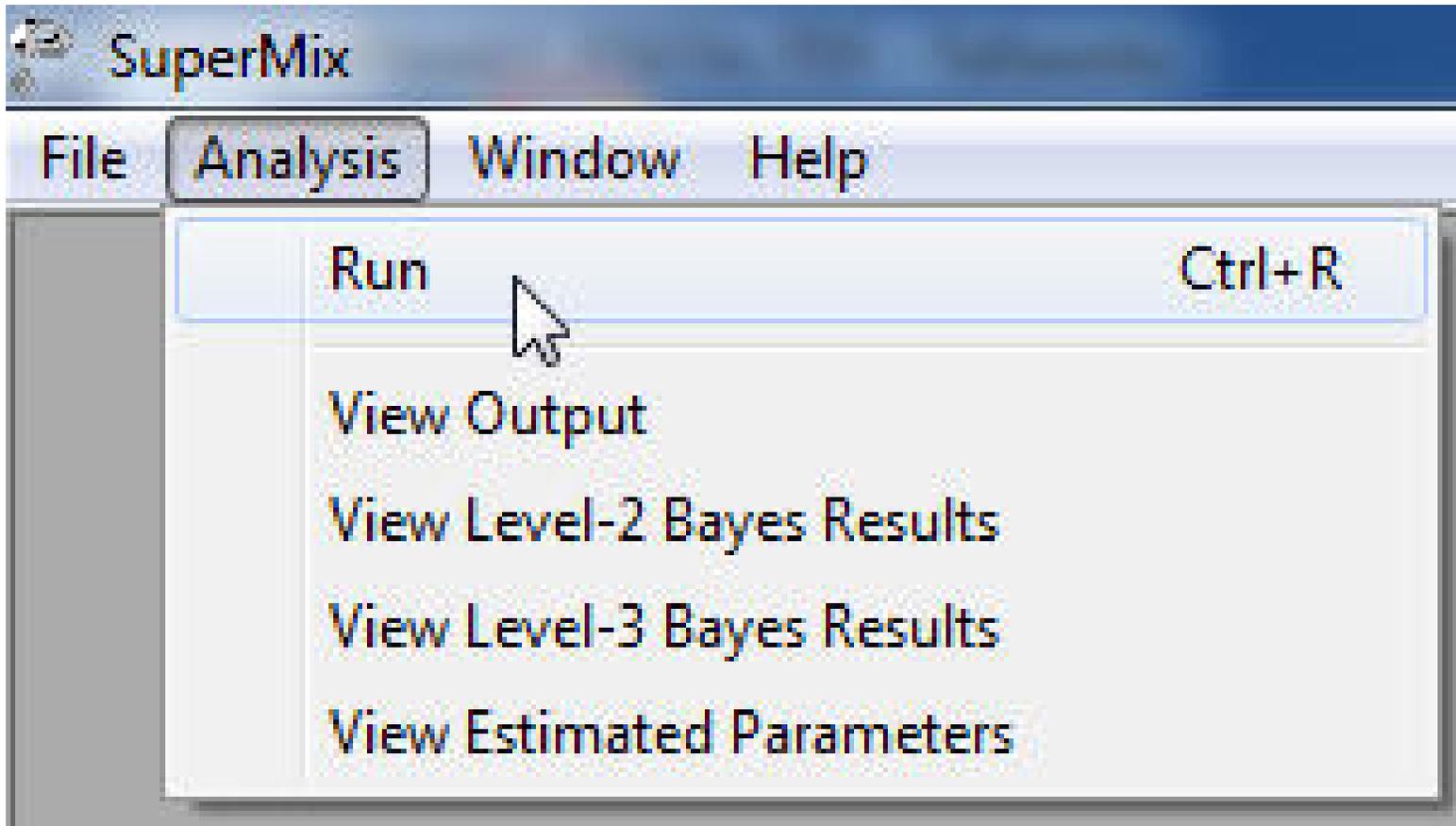
Dependent (Binary) Variable Settings

Distribution Model: Bernoulli

Function Model: logistic

Estimate Scale: none

Select the optimization method.
The default is adaptive quadrature.



schizb1.out

```

| Schiz BINARY outcome |
| random intercept model |
|-----|

```

Model and Data Descriptions

```

Sampling Distribution           = Bernoulli
Link Function                  = Logistic
PROB(Success)= 1.0/[1.0+EXP(-ETA)]

```

```

Number of Level-2 Units           437
Number of Level-1 Units          1603
Number of Level-1 Units per Level-2 Unit =

```

4	4	3	4	4	4	4	4	4	3	4	4
4	2	3	4	3	4	3	4	4	4	3	3
2	4	4	4	4	4	3	4	4	4	4	4
4	4	4	4	2	3	4	3	4	4	4	3
4	4	2	2	4	5	4	2	4	4	3	4
4	3	2	3	4	4	4	4	4	4	2	4
4	4	5	4	4	2	2	4	2	4	4	3
3	4	4	4	4	4	4	4	4	3	3	4
2	3	4	4	4	2	5	3	4	4	2	4
4	4	2	4	4	4	4	4	4	4	4	4
5	2	4	3	4	4	2	2	4	4	4	4
4	2	4	4	4	4	4	4	4	4	4	4
4	4	4	2	4	4	2	4	4	4	3	4
2	4	4	3	2	3	4	4	3	3	4	3
4	4	4	4	4	4	4	4	4	4	4	4
4	4	2	3	3	5	4	3	4	4	3	2
4	4	4	4	4	3	3	4	4	4	4	4
4	4	4	4	4	4	4	4	4	4	3	4
4	4	4	4	4	2	3	4	4	4	4	4
4	4	4	3	4	4	4	4	4	4	4	4
3	4	4	3	4	4	2	4	4	4	4	2
4	4	4	2	4	4	4	3	3	4	3	4
2	4	4	4	3	3	4	4	4	4	3	3
4	3	4	4	4	4	4	3	4	4	4	4
4	3	3	4	2	4	4	4	4	4	4	4
4	3	4	4	3	3	4	2	4	3	3	3
4	4	4	4	4	4	4	3	2	3	4	4
.

Save As... Close

schizbl.out

Optimization Method: Adaptive Quadrature

Number of quadrature points = 25
 Number of free parameters = 5
 Number of iterations used = 6

-2lnL (deviance statistic) = 1249.73465
 Akaike Information Criterion 1259.73465
 Schwarz Criterion 1286.63281

Estimated regression weights

Parameter	Estimate	Standard Error	z Value	P Value
intercept	5.3851	0.6303	8.5432	0.0000
TxDrug	-0.0247	0.6533	-0.0378	0.9698
SqrtWeek	-1.4996	0.2906	-5.1606	0.0000
Tx*SWeek	-1.0143	0.3338	-3.0385	0.0024

Odds Ratio and 95% Odds Ratio Confidence Intervals

Parameter	Estimate	Odds Ratio	Bounds	
			Lower	Upper
intercept	5.3851	218.1386	63.4125	750.3956
TxDrug	-0.0247	0.9756	0.2711	3.5106
SqrtWeek	-1.4996	0.2232	0.1263	0.3945
Tx*SWeek	-1.0143	0.3627	0.1885	0.6977

Estimated level 2 variances and covariances

Parameter	Estimate	Standard Error	z Value	P Value
intercept/intercept	4.4781	0.9458	4.7345	0.0000

Save As... Close

```

schizbl.out
-----
Calculation of the intraclass correlation
-----
residual variance = pi*pi / 3 (assumed)
cluster variance = 4.4781

intraclass correlation = 4.4781 / ( 4.4781 + (pi*pi/3)) = 0.576

Population Average Estimates

Parameter      Estimate      Standard
-----      -
intercept      3.5427       0.4628
TxDrug         -0.0546      0.5162
SqrtWeek       -1.0503      0.2238
Tx*SWeek       -0.5964      0.2502
z Value        P Value
-----        -
intercept      7.6549      0.0000
TxDrug         -0.1058     0.9157
SqrtWeek       -4.6936     0.0000
Tx*SWeek       -2.3838     0.0171

Odds Ratio and 95% Odds Ratio Confidence Intervals

Parameter      Estimate      Odds Ratio      Bounds
-----      -
intercept      3.5427       34.5605      Lower      Upper
TxDrug         -0.0546      0.9468      0.3443     2.6040
SqrtWeek       -1.0503      0.3498      0.2256     0.5424
Tx*SWeek       -0.5964      0.5508      0.3373     0.8994

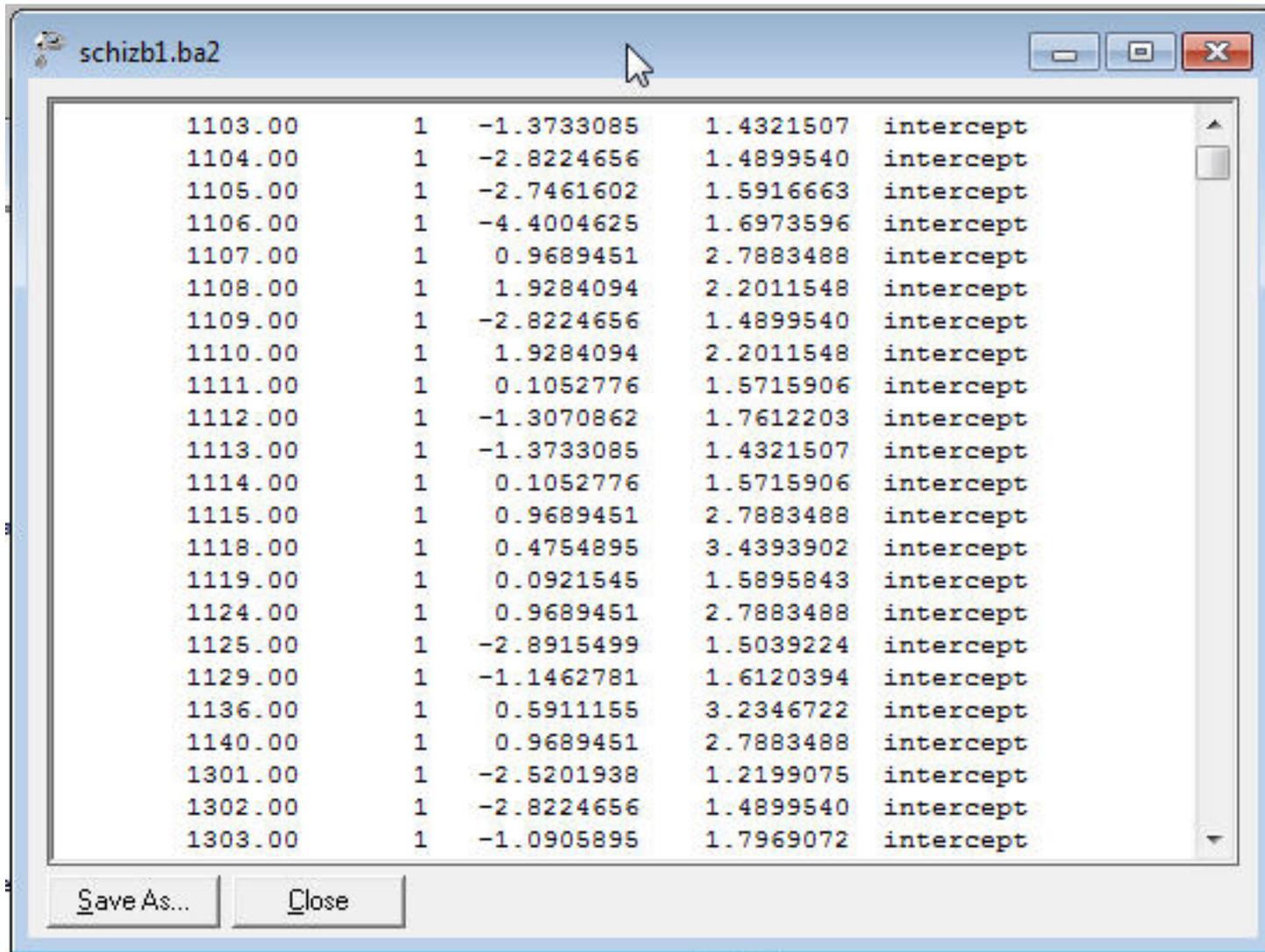
=====
| SuperMix used 0.31 seconds CPU |
=====

```

SuperMix is FAST for full-likelihood estimation up to 3-level models

Empirical Bayes Estimates of Random Effects

Select “Analysis” > “View Level-2 Bayes Results”



ID	random effect number	estimate	variance	name
1103.00	1	-1.3733085	1.4321507	intercept
1104.00	1	-2.8224656	1.4899540	intercept
1105.00	1	-2.7461602	1.5916663	intercept
1106.00	1	-4.4004625	1.6973596	intercept
1107.00	1	0.9689451	2.7883488	intercept
1108.00	1	1.9284094	2.2011548	intercept
1109.00	1	-2.8224656	1.4899540	intercept
1110.00	1	1.9284094	2.2011548	intercept
1111.00	1	0.1052776	1.5715906	intercept
1112.00	1	-1.3070862	1.7612203	intercept
1113.00	1	-1.3733085	1.4321507	intercept
1114.00	1	0.1052776	1.5715906	intercept
1115.00	1	0.9689451	2.7883488	intercept
1118.00	1	0.4754895	3.4393902	intercept
1119.00	1	0.0921545	1.5895843	intercept
1124.00	1	0.9689451	2.7883488	intercept
1125.00	1	-2.8915499	1.5039224	intercept
1129.00	1	-1.1462781	1.6120394	intercept
1136.00	1	0.5911155	3.2346722	intercept
1140.00	1	0.9689451	2.7883488	intercept
1301.00	1	-2.5201938	1.2199075	intercept
1302.00	1	-2.8224656	1.4899540	intercept
1303.00	1	-1.0905895	1.7969072	intercept

ID, random effect number, estimate, variance, name

Select “Model-based Graphs” > “Equations”

Plot Equations for Outcome Variable

List of Variables

Name	Predictor	Group	Mark
intercept	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
TxDrug	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SqrtWeek	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Tx*S/week	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Patient		<input type="checkbox"/>	<input checked="" type="checkbox"/>

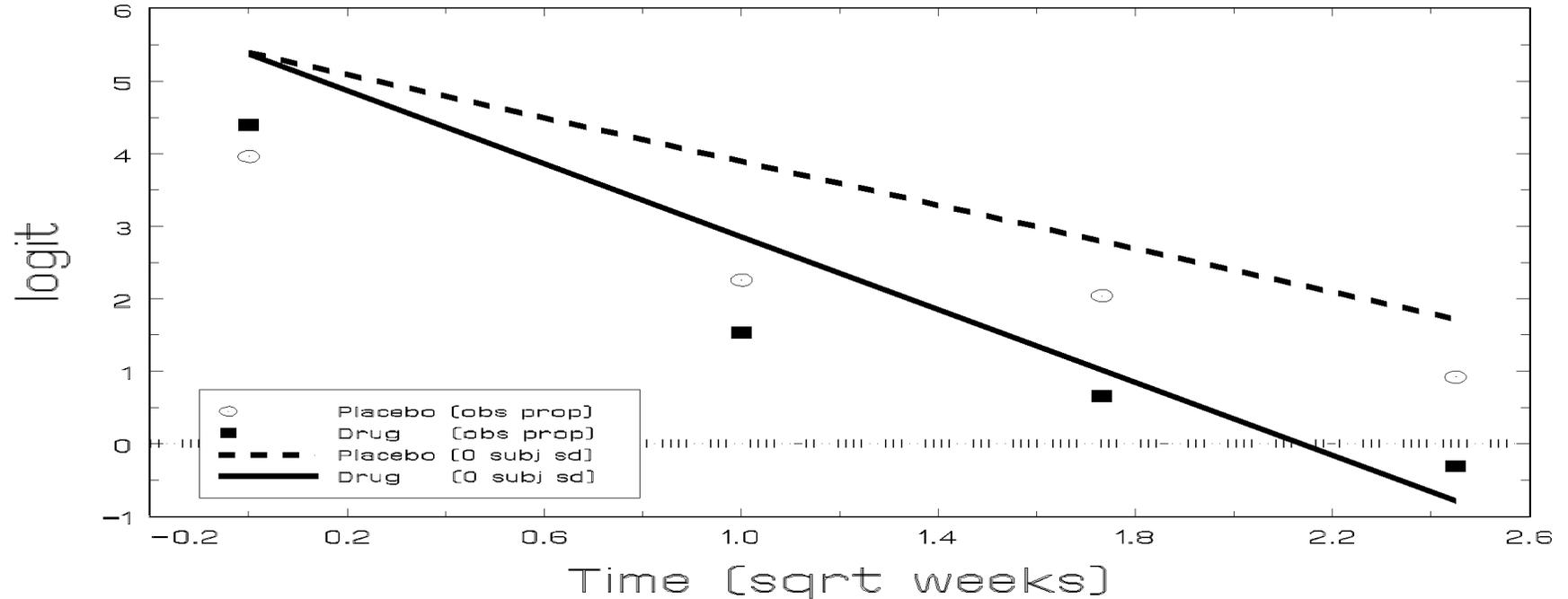
Remaining predictors fixed at 0
 Remaining predictors fixed at their means
 Plot linear regression model

Note: Only one variable may be selected for grouping and only one for marking.

Plot Cancel

Estimated (subject-specific) Logits across Time by Condition: *random-intercepts model*

Random Intercepts Logistic Model



$$\log \left[\frac{Pr(Y_{ij} = 1)}{1 - Pr(Y_{ij} = 1)} \right] = 5.39 - .03 D_i - 1.50 T_j - 1.01 (D_i \times T_j) + v_{0i}$$

$$v_{0i} \sim \mathcal{NID}(0, \hat{\sigma}_v^2 = 4.48)$$

$\hat{\beta}$ change in (conditional) logit due to \mathbf{x} for subjects with the same value of v_{0i} (the above plot is for $v_{0i} = 0$)

Random-intercepts Logistic Regression

$$\text{logit}_{ij} = \mathbf{x}'_{ij}\boldsymbol{\beta} + v_{0i}$$

- every subject has their own propensity for response (v_{0i})
- the influence of covariates \mathbf{x} is determined controlling (or adjusting) for the subject effect
- the covariance structure, or dependency, of the repeated observations is explicitly modeled

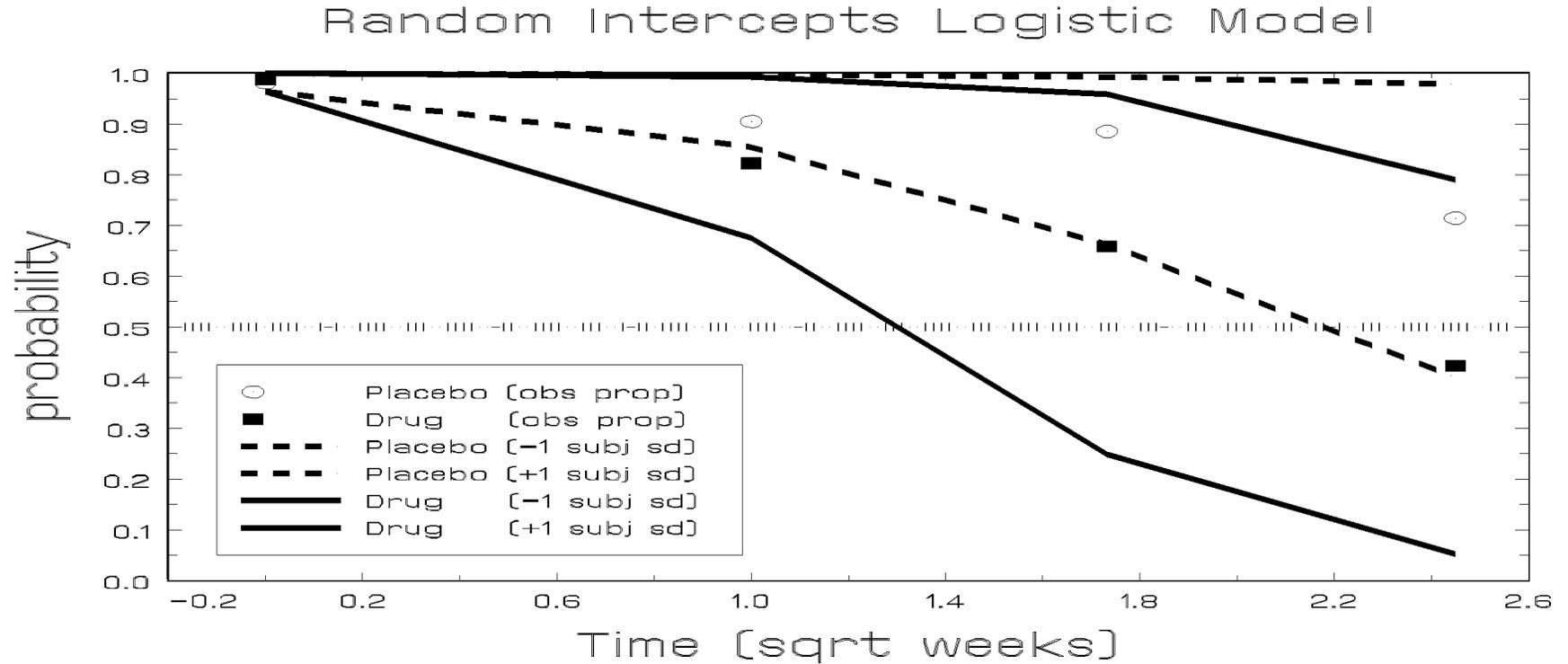
β_0 = log odds of response for a typical subject with $\mathbf{x} = 0$ and $v_{0i} = 0$

β = log odds ratio for response associated with unit changes in \mathbf{x} for the same subject value v_{0i}
* referred to as “subject-specific”
* how a *subject's* response probability depends on \mathbf{x}

σ_v^2 = degree of heterogeneity across subjects in the probability of response not attributable to \mathbf{x}

- most useful when the objective is to make inference about *subjects* rather than the population average
- interest is in the heterogeneity of subjects

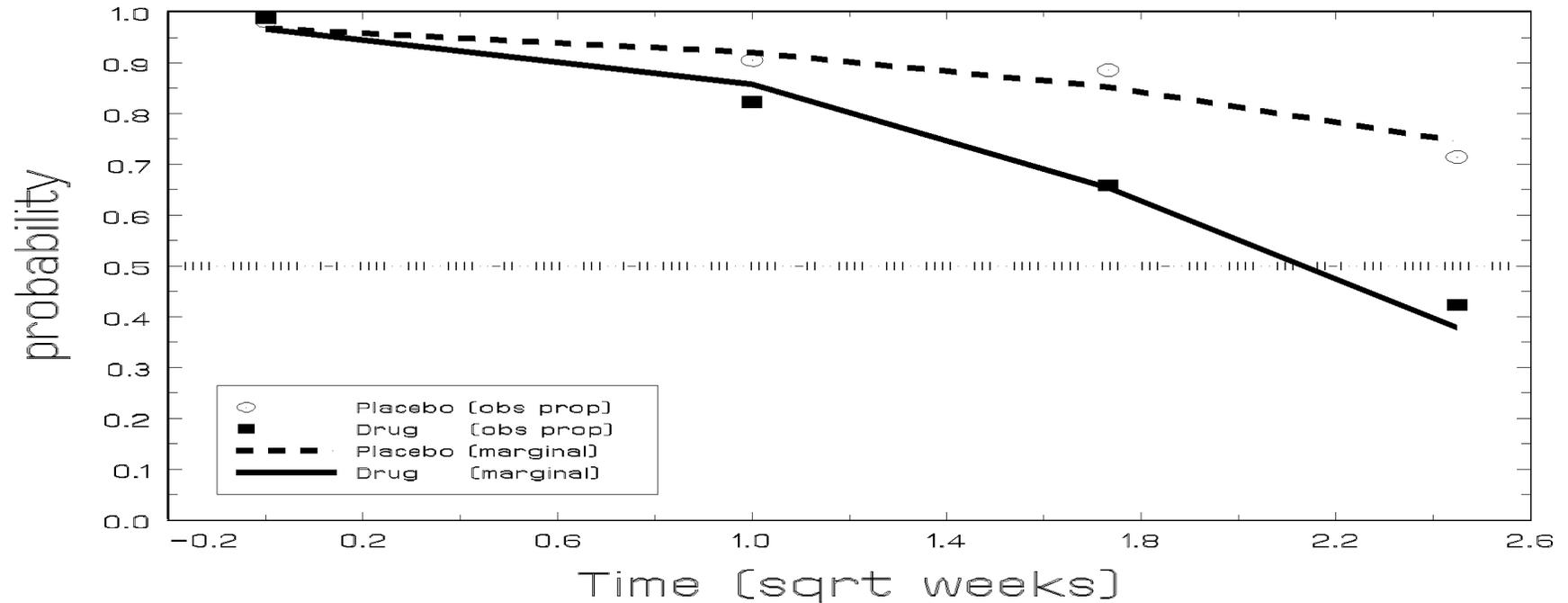
Estimated Subject-Specific Probabilities



$$Pr(Y_{ij} = 1) = \frac{1}{1 + \exp \left[- \left(5.39 - .03 D_i - 1.50 T_j - 1.01 D_i T_j + v_{0i} \right) \right]}$$

$$\text{where } v_{0i} = \begin{cases} -1\sigma_v \\ 1\sigma_v \end{cases} \text{ and } \hat{\sigma}_v = 2.12$$

Marginalized Random Intercepts Logistic

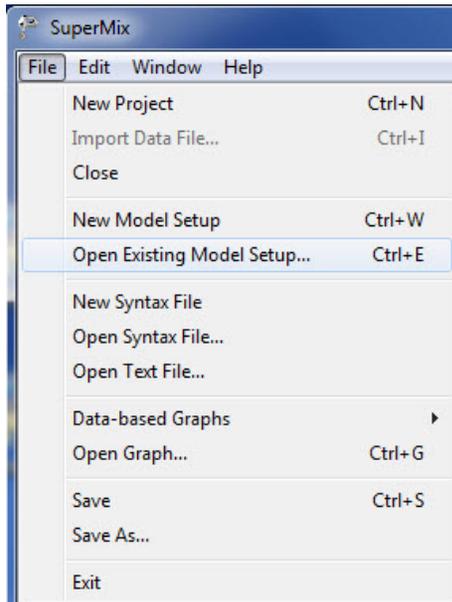


$$P(Y_{ij} = 1) = \frac{1}{1 + \exp \left[- \left(3.54 - .055 D_i - 1.05 T_j - .60 D_i T_j \right) \right]}$$

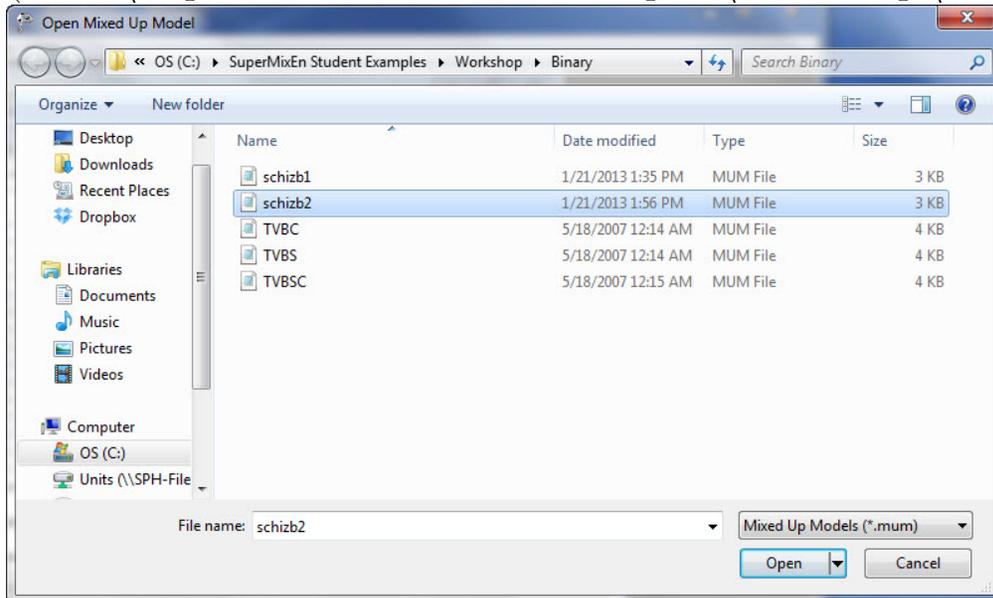
⇒ obtained using the Population Average estimates from Supermix

Hedeker D, du Toit SHC, Demirtas H, & Gibbons RD (2018). A note on marginalization of regression parameters from mixed models of binary outcomes. *Biometrics*, 74(1):354-361.

Under “File” click on “Open Existing Model Setup”



Open C:\SuperMixEn Examples\Workshop\Binary\schizb2.mum
(or C:\SuperMixEn Student Examples\Workshop\Binary\schizb2.mum)



Note “Dependent Variable Type” should be “binary”

Model Setup: schizb2.mum

Configuration | Variables | Starting Values | Patterns | Advanced | Linear Transforms

Title 1: Schiz BINARY outcome

Title 2: random intercept and trend model

Dependent Variable Type: **binary**

Level-2 IDs: Patient

Dependent Variable: Imps79D

Level-3 IDs:

Categories:

	Value
1	0
2	1

Write Bayes Estimates: no

Convergence Criterion: 0.001

Number of Iterations: 100

Missing Values Present: true

Perform Crosstabulation: no

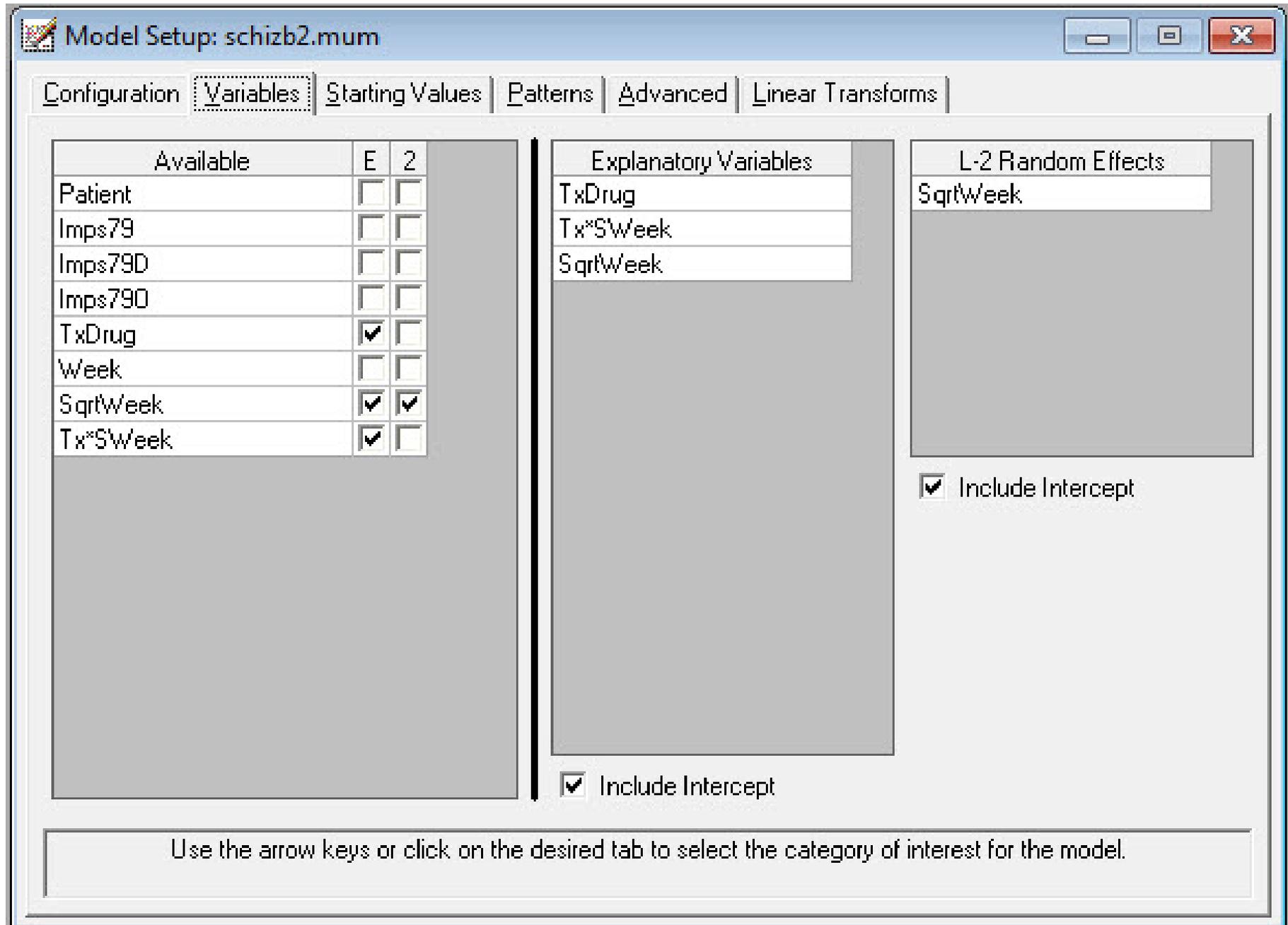
Missing Value for the Dependent Var: -9.0

Global Missing Value: -9.0

Output Type: standard

Select the form of the dependent variable. The options on the screens will change as required.

SqrtWeek is a level-2 (subject) random effect



Model Setup: schizb2.mum

Configuration Variables Starting Values Patterns Advanced Linear Transforms

Available	E	2
Patient	<input type="checkbox"/>	<input type="checkbox"/>
Imps79	<input type="checkbox"/>	<input type="checkbox"/>
Imps79D	<input type="checkbox"/>	<input type="checkbox"/>
Imps790	<input type="checkbox"/>	<input type="checkbox"/>
TxDrug	<input checked="" type="checkbox"/>	<input type="checkbox"/>
Week	<input type="checkbox"/>	<input type="checkbox"/>
SqrtWeek	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
Tx*S'week	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Explanatory Variables

- TxDrug
- Tx*S'week
- SqrtWeek

Include Intercept

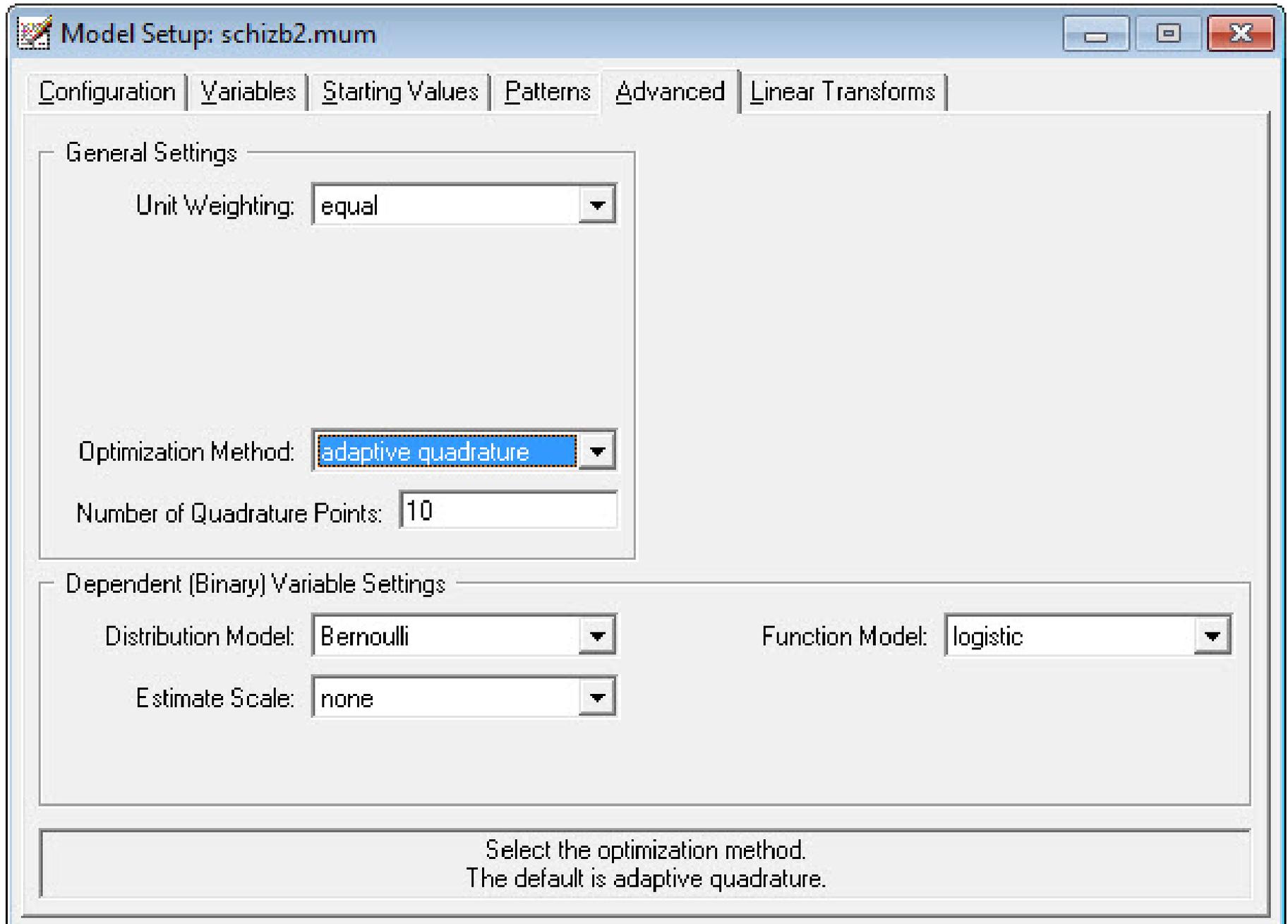
L-2 Random Effects

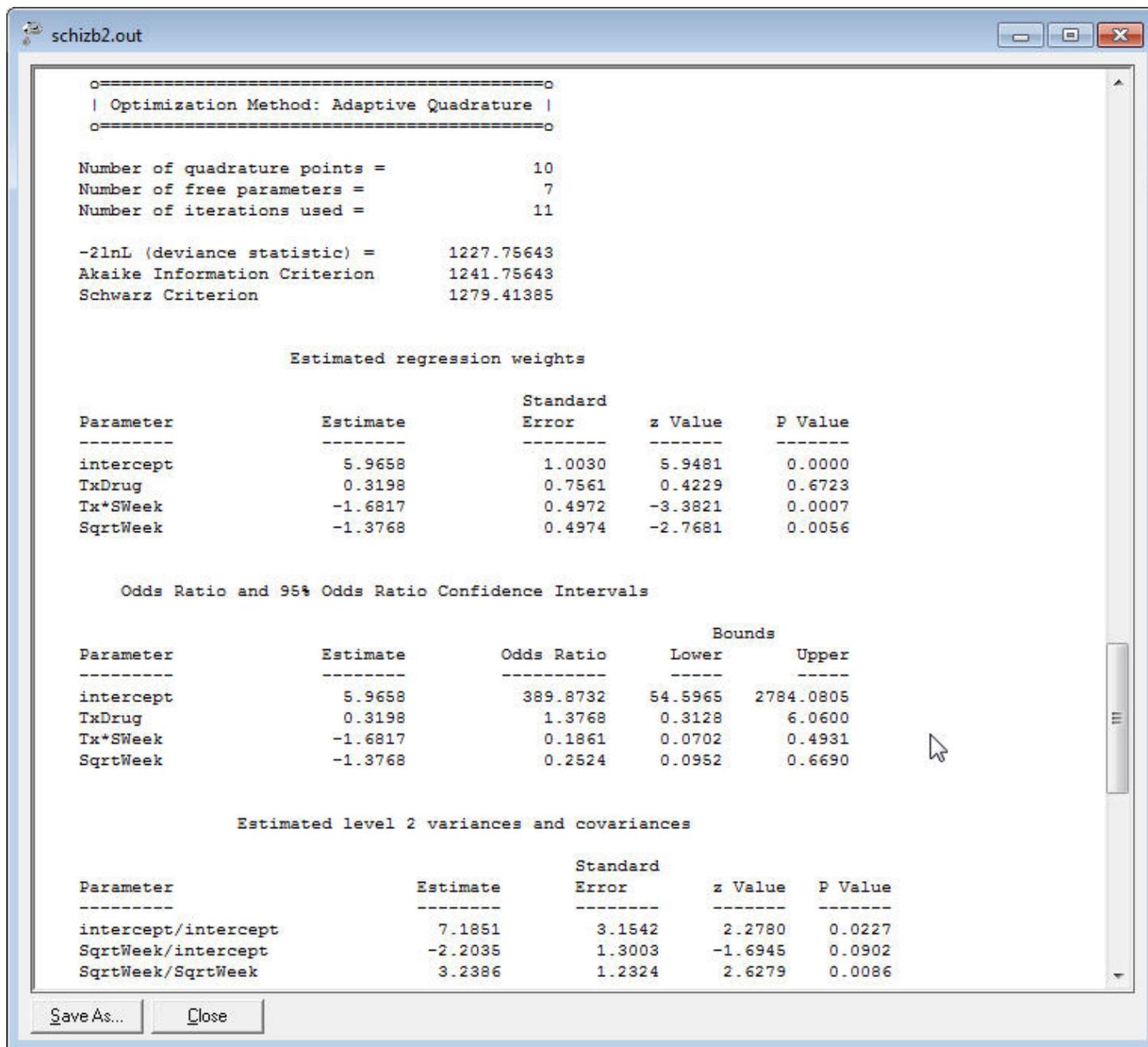
- SqrtWeek

Include Intercept

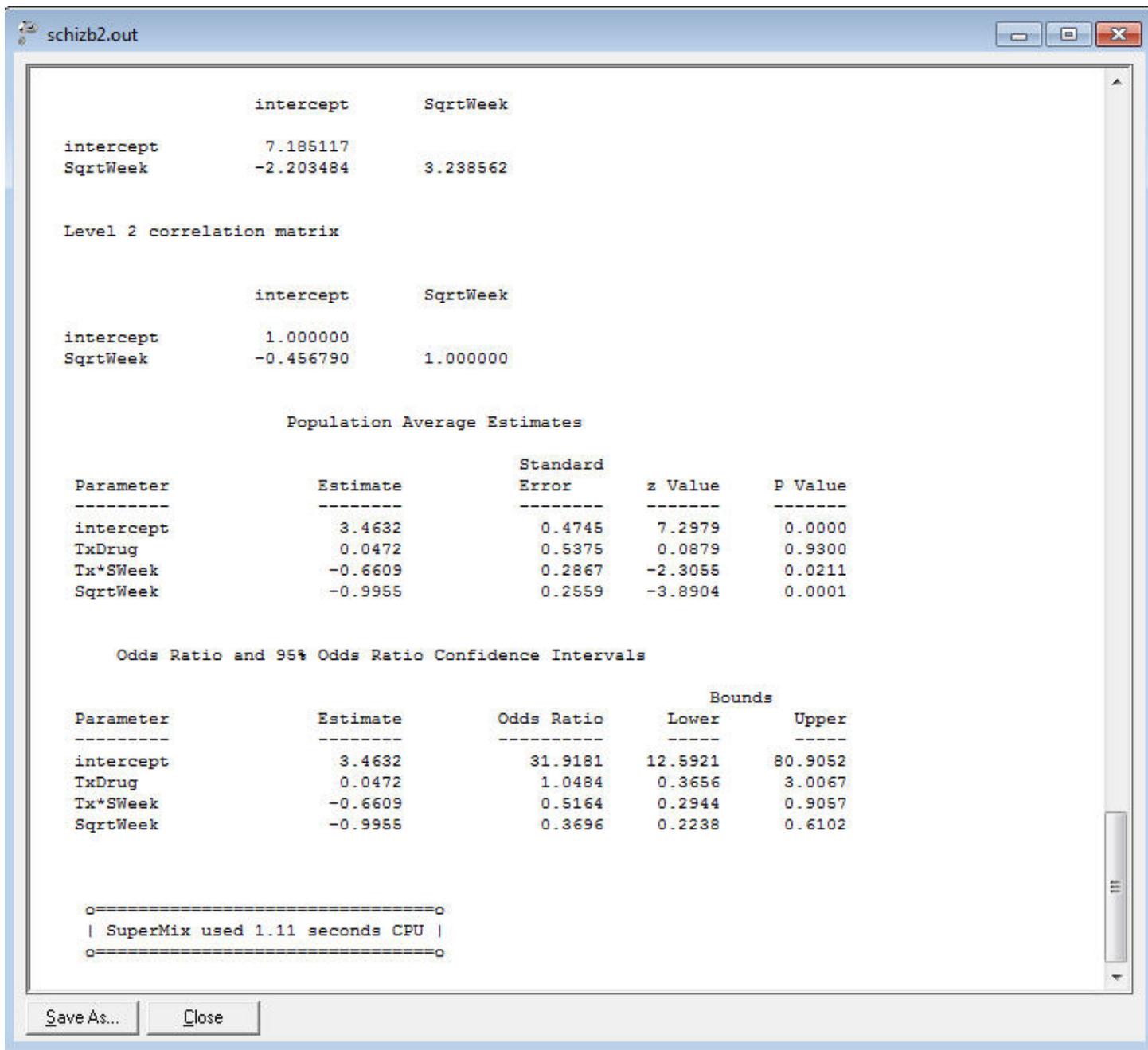
Use the arrow keys or click on the desired tab to select the category of interest for the model.

Note “Optimization Method” should be “adaptive quadrature”





⇒ Comparing models: $H_0 : \sigma_{v_1}^2 = \sigma_{v_{01}} = 0$; $\chi_2^2 = 1249.73 - 1227.76 = 21.97, p < .001$



Select “Model-based Graphs” > “Equations”

Plot Equations for Outcome Variable

List of Variables

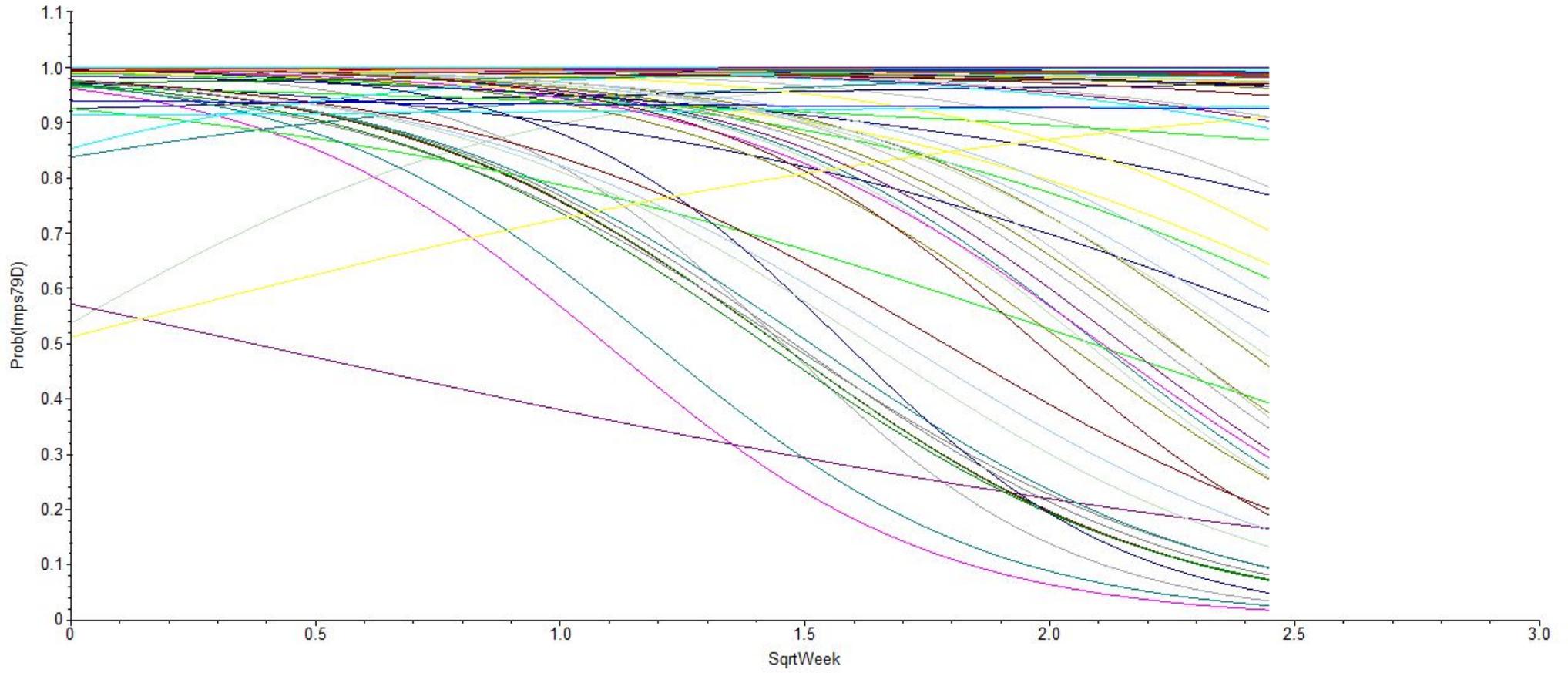
Name	Predictor	Group	Mark
intercept	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
TxDrug	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
SqrtWeek	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Tx*S/week	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>
Patient		<input type="checkbox"/>	<input checked="" type="checkbox"/>

Remaining predictors fixed at 0
 Remaining predictors fixed at their means
 Plot linear regression model

Note: Only one variable may be selected for grouping and only one for marking.

Plot Cancel

Prob(Imps79D) vs. SqrtWeek



Supermix - mixed models for binary outcomes

- link functions: logistic, probit, log-log, complementary log-log
- multiple random effects (correlated or independent) for up to 3-level models
- fast full-likelihood estimation using adaptive Gauss-Hermite quadrature
- yields both subject-specific (conditional) and population-average (marginal) estimates and inference of regression coefficients
- discrete/grouped time survival analysis via person-period dataset
www.ssicentral.com/images/pdfs/Survival_clust.pdf