

## The SAS System

### The GLIMMIX Procedure

Model Information	
Data Set	WORK.Y06_2YR
Response Variable (Events)	dmg_now
Response Variable (Trials)	tot_nuts
Response Distribution	Binomial
Link Function	Logit
Variance Function	Default
Variance Matrix	Not blocked
Estimation Technique	Residual PL
Degrees of Freedom Method	Kenward-Roger
Fixed Effects SE Adjustment	Kenward-Roger

Class Level Information		
Class	Levels	Values
Year	2	2006 2007
Tier	8	1 2 3 4 7 8 9 10
treatment	3	C CMD MD

Number of Observations Read	16
Number of Observations Used	16
Number of Events	1347
Number of Trials	135524

Dimensions	
G-side Cov. Parameters	3
Columns in X	4
Columns in Z	26
Subjects (Blocks in V)	1
Max Obs per Subject	16

Optimization Information	
Optimization Technique	Dual Quasi-Newton
Parameters in Optimization	3
Lower Boundaries	3

<b>Upper Boundaries</b>	0
<b>Fixed Effects</b>	Profiled
<b>Starting From</b>	Data

<b>Iteration History</b>					
<b>Iteration</b>	<b>Restarts</b>	<b>Subiterations</b>	<b>Objective Function</b>	<b>Change</b>	<b>Max Gradient</b>
0	0	5	22.039864332	0.01368619	4.158E-6
1	0	4	22.05249808	0.00060503	7.119E-8
2	0	2	22.052606981	0.00000513	6.014E-7
3	0	0	22.052607587	0.00000000	1.499E-6

Convergence criterion (PCONV=1.11022E-8) satisfied.

<b>Fit Statistics</b>	
<b>-2 Res Log Pseudo-Likelihood</b>	22.05
<b>Generalized Chi-Square</b>	12.90
<b>Gener. Chi-Square / DF</b>	0.99

<b>Covariance Parameter Estimates</b>		
<b>Cov Parm</b>	<b>Estimate</b>	<b>Standard Error</b>
<b>Year</b>	0.5263	0.7687
<b>Tier</b>	0.06173	0.08582
<b>Year*Tier</b>	0.1070	0.06719

<b>Type III Tests of Fixed Effects</b>				
<b>Effect</b>	<b>Num DF</b>	<b>Den DF</b>	<b>F Value</b>	<b>Pr &gt; F</b>
treatment	2	7.052	6.59	0.0243

<b>treatment Least Squares Means</b>					
<b>treatment</b>	<b>Estimate</b>	<b>Standard Error</b>	<b>DF</b>	<b>t Value</b>	<b>Pr &gt;  t </b>
C	-4.6976	0.5401	1.136	-8.70	0.0560
CMD	-5.9503	0.6027	1.609	-9.87	0.0198
MD	-4.6370	0.5698	1.411	-8.14	0.0372

<b>Differences of treatment Least Squares Means Adjustment for Multiple Comparisons: Tukey-Kramer</b>					

treatment	_treatment	Estimate	Standard Error	DF	t Value	Pr >  t	Adj P
C	CMD	1.2526	0.3642	11.4	3.44	0.0053	0.0134
C	MD	-0.06065	0.3006	5.046	-0.20	0.8480	0.9779
CMD	MD	-1.3133	0.4010	7.376	-3.27	0.0126	0.0301

**Tukey-Kramer Grouping  
for treatment Least  
Squares Means (Alpha=0.05)**

**LS-means with the same  
letter are not significantly  
different.**

treatment	Estimate	
MD	-4.6370	A
		A
C	-4.6976	A
CMD	-5.9503	B