The SAS System

The GENMOD Procedure

Model Information						
Data Set	WORK.GOOGLE_MEDIAN					
Distribution	Poisson					
Link Function	Log					
Dependent Variable	Rating					

Number of Observations Read	10840
Number of Observations Used	10840

Class Level Information							
Class	Levels	Values					
Rating	39	1.33526419 0.470628434 1.163463261 1.224069506 1.253048642 1.547775252 0.0009995003 0.1831545431 0.3371862674 0.4061315527 0.5312163134 0.5883420662 0.6423800635 0.6936470556 0.7424134219 0.7889118025 0.8333438111 0.8758853172 0.9166906519 0.9558959865					
Free	2	01					

Criteria For Assessing Goodness Of Fit							
Criterion	DF	Value	Value/DF				
Deviance	11E3	175.1571	0.0162				
Scaled Deviance	11E3	175.1571	0.0162				
Pearson Chi-Square	11E3	139.4881	0.0129				
Scaled Pearson X2	11E3	139.4881	0.0129				
Log Likelihood		-9957.0611					
Full Log Likelihood		-12572.1057					
AIC (smaller is better)		25152.2114					
AICC (smaller is better)		25152.2151					
BIC (smaller is better)		25181.3754					

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates										
Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square	Pr > ChiSq		
Intercept		1	0.4115	0.0329	0.3471	0.4759	156.89	<.0001		
Installs		1	-0.0198	0.0070	-0.0335	-0.0062	8.08	0.0045		
Reviews		1	0.0228	0.0071	0.0089	0.0367	10.30	0.0013		
Free	0	1	-0.0080	0.0327	-0.0720	0.0560	0.06	0.8061		

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates									
Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square	Pr > ChiSq	
Free	1	0	0.0000	0.0000	0.0000	0.0000			
Scale		0	1.0000	0.0000	1.0000	1.0000			

Note: The scale parameter was held fixed.