The SURVEYFREQ Procedure

Data Summary	
Number of Observations	1447
Sum of Weights	4363

	Table of hosp by hosp_num						
hosp	hosp_num	Frequency	Weighted Frequency	Std Err of Wgt Freq	Percent	Std Err of Percent	
No	0	735	2808	75.19838	64.3594	1.3469	
	1	0					
	Total	735	2808	75.19838	64.3594	1.3469	
Yes	0	0					
	1	712	1555	57.08056	35.6406	1.3469	
	Total	712	1555	57.08056	35.6406	1.3469	
Total	0	735	2808	75.19838	64.3594	1.3469	
	1	712	1555	57.08056	35.6406	1.3469	
	Total	1447	4363	53.60645	100.0000		

Table of co						
со	Frequency	Weighted Frequency	Std Err of Wgt Freq	Percent	Std Err of Percent	
Community-onset	846	2907	72.90100	66.6285	1.3382	
Healthcare-associated	601	1456	58.42348	33.3715	1.3382	
Total	1447	4363	53.60645	100.0000		

The FREQ Procedure

hos	sp	hosp_num	Frequency	Percent	Cumulative Frequency	Cumulative Percent
No		0	2808	64.36	2808	64.36
Yes	;	1	1555	35.64	4363	100.00

со	Frequency	Percent	Cumulative Frequency	Cumulative Percent
Community-onset	2907	66.63	2907	66.63
Healthcare-associated	1456	33.37	4363	100.00

The FREQ Procedure

со	Frequency	Percent	Cumulative Frequency	Cumulative Percent
Community-onset	2907	100.00	2907	100.00

The FREQ Procedure

со	Frequency	Percent	Cumulative Frequency	Cumulative Percent
Healthcare-associated	1456	100.00	1456	100.00

The SURVEYLOGISTIC Procedure

Model Information			
Data Set	WORK.STAPH		
Response Variable	hosp		
Number of Response Levels	2		
Stratum Variable	INVASIVE		
Number of Strata	2		
Weight Variable	WEIGHT		
Model	Binary Logit		
Optimization Technique	Newton-Raphson		
Variance Adjustment	Degrees of Freedom (DF)		

Variance Estimation			
Method	Taylor Series		
Variance Adjustment Degrees of Freedom (DF			

Number of Observations Read	1447
Number of Observations Used	1447
Sum of Weights Read	4363
Sum of Weights Used	4363

Response Profile					
Ordered Value	hosp	Total Frequency	Total Weight		
1	No	735	2808.0000		
2	Yes	712	1555.0000		

Probability modeled is hosp='Yes'.

Class Level Information			
Value	Design Variables		
No	0		
Yes	1		
MRSA	1		
MSSA	0		
No	0		
Yes	1		
No	0		
Yes	1		
	Value No Yes MRSA MSSA No Yes No		

The SURVEYLOGISTIC Procedure

Class Level Information					
Class	Class Value Variables				
SMOKER	No	0			
	Yes	1			
BSI	No	0			
	Yes	1			

Stratum Information				
Stratum INVASIVE N Obs				
1	No	953		
2	Yes	494		

Model Convergence Status

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

Model Fit Statistics					
Intercept and Criterion Only Covariates					
AIC	5685.439	4134.144			
sc	5691.820	4185.191			
-2 Log L	5683.439	4118.144			

	R-Square	0.3015	Max-rescaled R-Square	0.4140	ı
-1					ł

Testing Global Null Hypothesis: BETA=0						
Test	F Value	Num DF	Den DF	Pr > F		
Likelihood Ratio	74.16	7.0000	10115	<.0001		
Score	121.72	7	1439	<.0001		
Wald	39.21	7	1439	<.0001		

NOTE: Second-order Rao-Scott design correction 0.0000 applied to the Likelihood Ratio test.

The SURVEYLOGISTIC Procedure

Type 3 Analysis of Effects							
Effect F Value Num DF Den DF Pr							
SMOKER	22.54	1	1445	<.0001			
mrsafinal	22.41	1	1445	<.0001			
kidney	13.38	1	1445	0.0003			
DIABETES	35.62	1	1445	<.0001			
BSI	158.48	1	1445	<.0001			
WOUND	15.02	1	1445	0.0001			
newage	9.69	1	1445	0.0019			

Analysis of Maximum Likelihood Estimates					
Parameter		Estimate	Standard Error	t Value	Pr > t
Intercept		-2.3264	0.1864	-12.48	<.0001
SMOKER	Yes	1.0095	0.2126	4.75	<.0001
mrsafinal	MRSA	0.7667	0.1620	4.73	<.0001
kidney	Yes	1.0379	0.2838	3.66	0.0003
DIABETES	Yes	1.1657	0.1953	5.97	<.0001
BSI	Yes	3.6303	0.2884	12.59	<.0001
WOUND	Yes	0.6579	0.1697	3.88	0.0001
newage		0.0108	0.00346	3.11	0.0019
NOTE: 1	The dear	es of freed	om for the t	tests is 1	445.

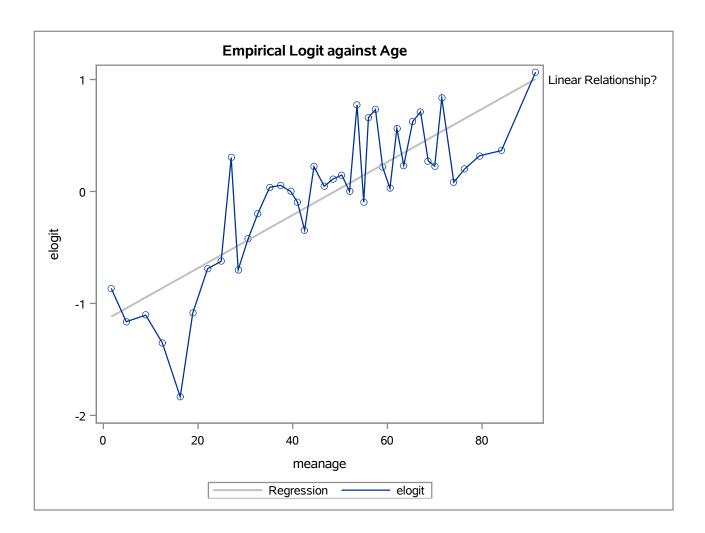
Odds Ratio Estimates					
Effect	Point Estimate				
SMOKER Yes vs No	2.744	1.808	4.164		
mrsafinal MRSA vs MSSA	2.153	1.567	2.958		
kidney Yes vs No	2.823	1.618	4.926		
DIABETES Yes vs No	3.208	2.187	4.706		
BSI Yes vs No	37.725	21.427	66.420		
WOUND Yes vs No	1.931	1.384	2.694		
newage	1.011	1.004	1.018		

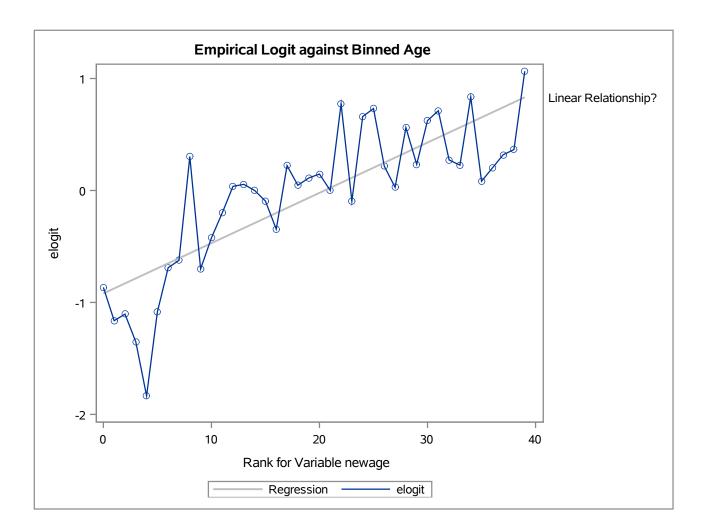
NOTE: The degrees of freedom in computing the confidence limits is 1445.

The SURVEYLOGISTIC Procedure

Association of Predicted Probabilities and Observed Responses					
Percent Concordant 86.4 Somers' D 0.731					
Percent Discordant	13.4	Gamma	0.732		
Percent Tied	0.2	Tau-a	0.366		
Pairs	523320	С	0.865		

	Estimated Correlation Matrix							
Parameter	Intercept	SMOKERYes	mrsafinalMRSA	kidneyYes	DIABETESYes	BSIYes	WOUNDYes	newage
Intercept	1.0000	-0.1785	-0.3593	0.0276	0.0005	-0.1332	-0.1622	-0.7818
SMOKERYes	-0.1785	1.0000	-0.0766	0.0801	0.0184	0.0226	-0.0322	0.0285
mrsafinalMRSA	-0.3593	-0.0766	1.0000	-0.0583	-0.0496	0.0480	0.0557	0.1120
kidneyYes	0.0276	0.0801	-0.0583	1.0000	-0.1249	-0.0402	0.0667	-0.1625
DIABETESYes	0.0005	0.0184	-0.0496	-0.1249	1.0000	0.1465	-0.0995	-0.1909
BSIYes	-0.1332	0.0226	0.0480	-0.0402	0.1465	1.0000	0.1665	-0.0821
WOUNDYes	-0.1622	-0.0322	0.0557	0.0667	-0.0995	0.1665	1.0000	-0.1344
newage	-0.7818	0.0285	0.1120	-0.1625	-0.1909	-0.0821	-0.1344	1.0000





The CORR Procedure

2 Variables: elogit meanage

Simple Statistics						
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum
elogit	40	-0.04369	0.65292	-1.74763	-1.83438	1.06538
meanage	40	47.02262	22.85461	1881	1.65789	91.29730

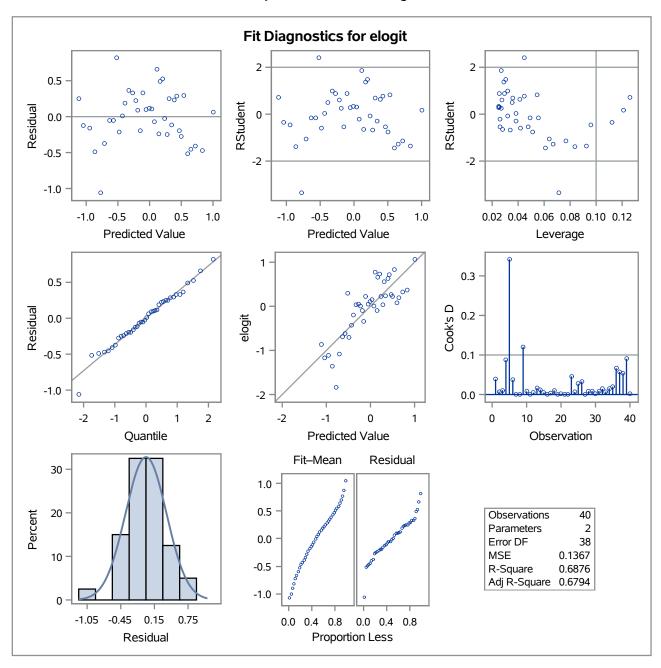
Pearson Correlation Coefficients, N = 40 Prob > r under H0: Rho=0					
elogit meanag					
elogit	1.00000	0.82924 <.0001			
meanage	0.82924 <.0001	1.00000			

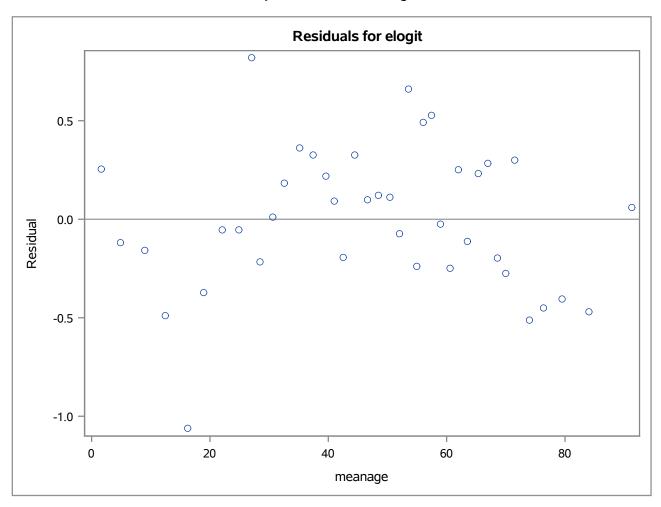
Number of Observations Read	40
Number of Observations Used	40

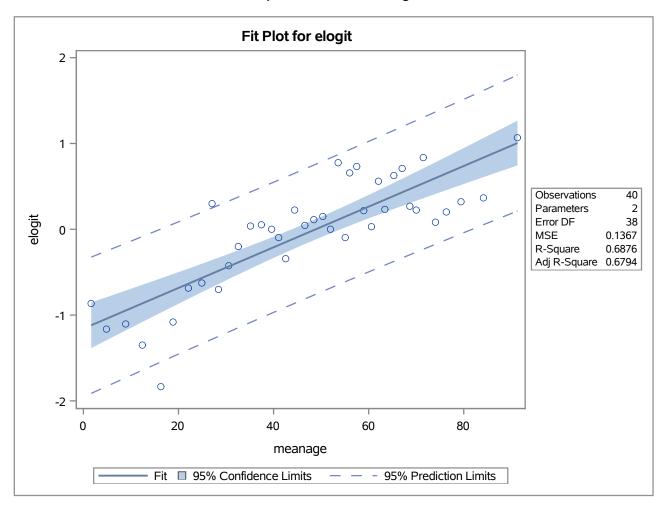
Analysis of Variance						
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F	
Model	1	11.43252	11.43252	83.66	<.0001	
Error	38	5.19311	0.13666			
Corrected Total	39	16.62563				

Root MSE	0.36968	R-Square	0.6876
Dependent Mean	-0.04369	Adj R-Sq	0.6794
Coeff Var	-846.12217		

Parameter Estimates						
Variable	DF	Parameter Estimate	Standard Error	t Value	Pr > t	
Intercept	1	-1.15766	0.13509	-8.57	<.0001	
meanage	1	0.02369	0.00259	9.15	<.0001	







The CORR Procedure

2 Variables: elogit bin

Simple Statistics							
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum	Label
elogit	40	-0.04369	0.65292	-1.74763	-1.83438	1.06538	
bin	40	19.50000	11.69045	780.00000	0	39.00000	Rank for Variable newage

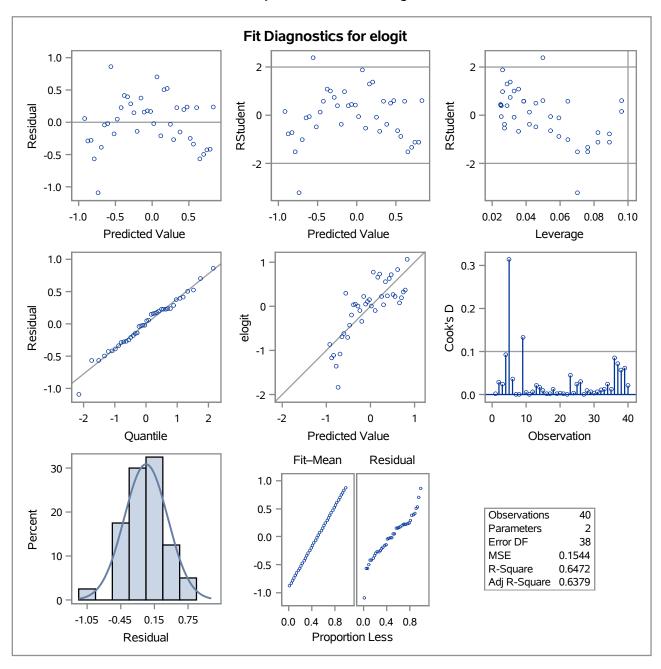
Pearson Correlation Coefficients, N = 40 Prob > r under H0: Rho=0					
	elogit	bin			
elogit	1.00000	0.80448 <.0001			
bin Rank for Variable newage	0.80448 <.0001	1.00000			

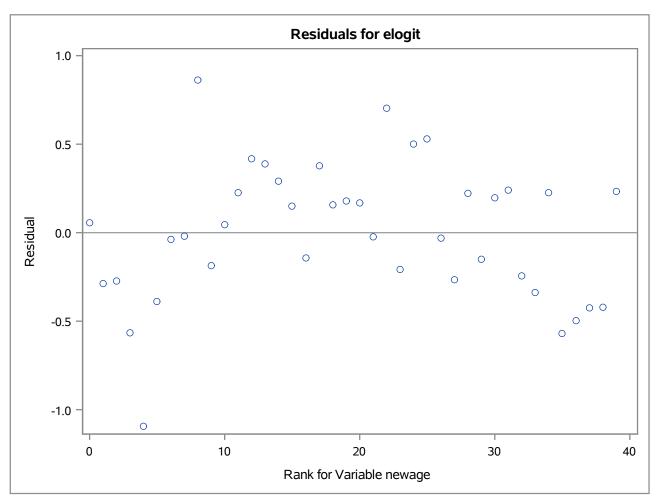
Number of Observations Read	40
Number of Observations Used	40

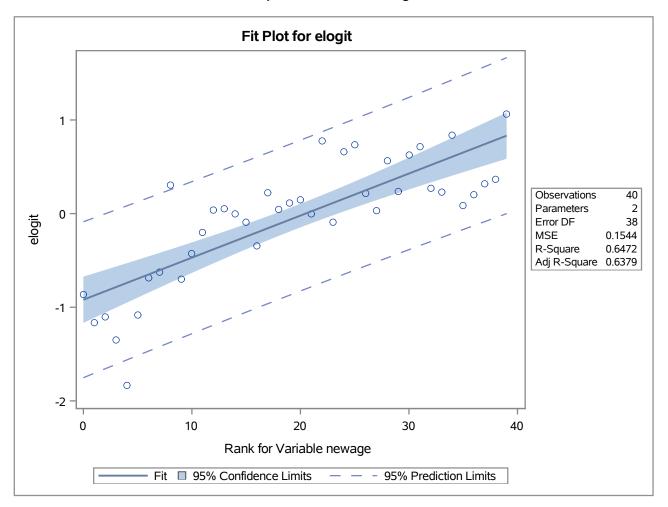
Analysis of Variance						
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F	
Model	1	10.76003	10.76003	69.71	<.0001	
Error	38	5.86560	0.15436			
Corrected Total	39	16.62563				

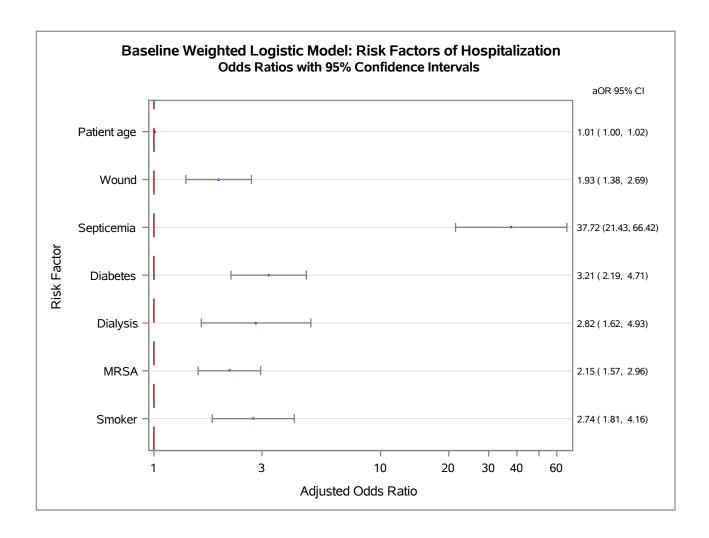
Root MSE	0.39288	R-Square	0.6472
Dependent Mean	-0.04369	Adj R-Sq	0.6379
Coeff Var	-899.23986		

Parameter Estimates						
Variable	Label	DF	Parameter Estimate	Standard Error	t Value	Pr > t
Intercept	Intercept	1	-0.91984	0.12195	-7.54	<.0001
bin	Rank for Variable newage	1	0.04493	0.00538	8.35	<.0001









Model Information			
Data Set	WORK.COMM		
Response Variable	hosp		
Number of Response Levels	2		
Stratum Variable	INVASIVE		
Number of Strata	2		
Weight Variable	WEIGHT		
Model	Binary Logit		
Optimization Technique	Newton-Raphson		
Variance Adjustment	Degrees of Freedom (DF)		

Variance Estimation				
Method	Taylor Series			
Variance Adjustment	Degrees of Freedom (DF)			

Number of Observations Read	846
Number of Observations Used	846
Sum of Weights Read	2907
Sum of Weights Used	2907

Response Profile				
Ordered Value	hosp	Total Frequency	Total Weight	
1	No	590	2285.0000	
2	Yes	256	622.0000	

Probability modeled is hosp='Yes'.

Class Level Information				
Value	Design Variables			
MRSA	1			
MSSA	0			
No	0			
Yes	1			
No	0			
Yes	1			
No	0			
Yes	1			
	Value MRSA MSSA No Yes No Yes No			

Class Level Information				
Class Value Design				
BSI	No	0		
	Yes	1		
WOUND	No	0		
	Yes	1		

Stratum Information				
Stratum INVASIVE N Obs				
1	No	679		
2	Yes	167		

Model Convergence Status

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

Model Fit Statistics				
Criterion	Intercept an Only Covaria			
AIC	3020.422	2305.551		
sc	3026.397	2353.350		
-2 Log L	3018.422	2289.551		

R-Square	0.2218	Max-rescaled R-Square	0.3433	
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Testing Global Null Hypothesis: BETA=0						
Test F Value Num DF Den DF Pr > F						
Likelihood Ratio	30.28	7.0000	5907.98	<.0001		
Score	36.84	7	838	<.0001		
Wald	24.99	7	838	<.0001		

NOTE:

Second-order Rao-Scott design correction 0.0000 applied to the Likelihood Ratio test.

Type 3 Analysis of Effects							
Effect	F Value Num DF Den DF Pr > F						
SMOKER	20.63	1	844	<.0001			
mrsafinal	9.00	1	844	0.0028			
kidney	2.37	1	844	0.1239			
DIABETES	18.60	1	844	<.0001			
BSI	123.15	1	844	<.0001			
WOUND	8.27	1	844	0.0041			
newage	0.65	1	844	0.4201			

Analysis of Maximum Likelihood Estimates					
Parameter		Estimate	Standard Error	t Value	Pr > t
Intercept		-2.5522	0.2434	-10.49	<.0001
SMOKER	Yes	1.2164	0.2678	4.54	<.0001
mrsafinal	MRSA	0.6836	0.2279	3.00	0.0028
kidney	Yes	0.7472	0.4851	1.54	0.1239
DIABETES	Yes	1.2194	0.2828	4.31	<.0001
BSI	Yes	4.5960	0.4142	11.10	<.0001
WOUND	Yes	0.6934	0.2410	2.88	0.0041
newage		0.00406	0.00504	0.81	0.4201
NOTE:	The degr	ees of freed	dom for the	t tests is 8	344.

Odds Ratio Estimates				
Effect	Point Estimate	95% Confidence Limits		
SMOKER Yes vs No	3.375	1.995 5.709		
mrsafinal MRSA vs MSSA	1.981	1.266	3.098	
kidney Yes vs No	2.111	0.815 5.471		
DIABETES Yes vs No	3.385	1.943 5.897		
BSI Yes vs No	99.091	43.954	223.391	
WOUND Yes vs No	2.000	1.246	3.211	
newage	1.004	0.994	1.014	

NOTE:

The degrees of freedom in computing the confidence limits is 844.

Association of Predicted Probabilities and Observed Responses						
Percent Concordant 83.3 Somers' D 0.674						
Percent Discordant 15.9 Gamma 0.67						
Percent Tied 0.8 Tau-a 0.28						
Pairs	151040	С	0.837			

Model Information			
Data Set WORK.HOP			
Response Variable	hosp		
Number of Response Levels	2		
Stratum Variable	INVASIVE		
Number of Strata	2		
Weight Variable	WEIGHT		
Model	Binary Logit		
Optimization Technique	Newton-Raphson		
Variance Adjustment	Degrees of Freedom (DF)		

Variance Estimation			
Method Taylor Series			
Variance Adjustment	Degrees of Freedom (DF)		

Number of Observations Read	601
Number of Observations Used	601
Sum of Weights Read	1456
Sum of Weights Used	1456

	Response Profile				
Ordered Total Total Value hosp Frequency Weight					
1	No	145	523.00000		
2	Yes	456	933.00000		

Probability modeled is hosp='Yes'.

Class Level Information			
Value	Design Variables		
MRSA	1		
MSSA	0		
No	0		
Yes	1		
No	0		
Yes	1		
No	0		
Yes	1		
	Value MRSA MSSA No Yes No Yes No		

Class Level Information			
Class	Value	Design Variables	
BSI	No	0	
	Yes	1	
WOUND	No	0	
	Yes	1	

Stratum Information			
Stratum Index	INVASIVE	N Obs	
1	No	274	
2	Yes	327	

Model Convergence Status

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

Model Fit Statistics				
Criterion	Intercept Only	Intercept and Covariates		
AIC	1903.415	1544.926		
sc	1908.698	1587.194		
-2 Log L	1901.415	1528.926		

R-Square	0.2257	Max-rescaled R-Square	0.3096
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Testing Global Null Hypothesis: BETA=0						
Test F Value Num DF Den DF Pr > F						
Likelihood Ratio	21.95	6.9999	4192.96	<.0001		
Score	593	<.0001				
Wald	10.58	7	593	<.0001		

NOTE:

Second-order Rao-Scott design correction 0.0000 applied to the Likelihood Ratio test.

Type 3 Analysis of Effects								
Effect	Effect F Value Num DF Den DF Pr > F							
SMOKER	4.50	1	599	0.0342				
mrsafinal	7.76	1	599	0.0055				
kidney	4.68	1	599	0.0309				
DIABETES	12.32	1	599	0.0005				
BSI	36.32	1	599	<.0001				
WOUND	0.17	1	599	0.6819				
newage	1.27	1	599	0.2608				

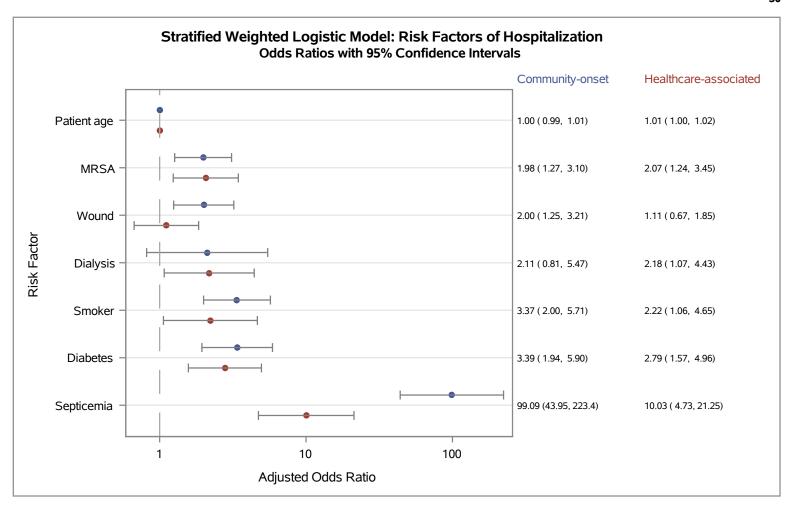
Analysis of Maximum Likelihood Estimates					
Parameter		Estimate	Standard Error	t Value	Pr > t
Intercept		-0.9211	0.3420	-2.69	0.0073
SMOKER	Yes	0.7982	0.3761	2.12	0.0342
mrsafinal	MRSA	0.7257	0.2606	2.79	0.0055
kidney	Yes	0.7799	0.3605	2.16	0.0309
DIABETES	Yes	1.0265	0.2924	3.51	0.0005
BSI	Yes	2.3051	0.3825	6.03	<.0001
WOUND	Yes	0.1059	0.2583	0.41	0.6819
newage		0.00669	0.00594	1.13	0.2608
NOTE: The degrees of freedom for the t tests is 599.					

Odds Ratio Estimates					
Effect	Point Estimate	95% Confidence Limits			
SMOKER Yes vs No	2.222	1.061	4.650		
mrsafinal MRSA vs MSSA	2.066	1.239	3.447		
kidney Yes vs No	2.181	1.075	4.428		
DIABETES Yes vs No	2.791	1.572	4.957		
BSI Yes vs No	10.025	4.730	21.248		
WOUND Yes vs No	1.112	0.669	1.846		
newage	1.007	0.995	1.019		

NOTE:

The degrees of freedom in computing the confidence limits is 599.

Association of Predicted Probabilities and Observed Responses						
Percent Concordant 81.3 Somers' D 0.630						
Percent Discordant	18.3	Gamma	0.632			
Percent Tied	0.4 Tau-a (0.231			
Pairs	66120	С	0.815			



The GLIMMIX Procedure

Model Information			
Data Set	S.STAPH		
Response Variable	hosp		
Response Distribution	Binary		
Link Function	Logit		
Variance Function	Default		
Variance Matrix Blocked By	TXHOSP		
Estimation Technique	Maximum Likelihood		
Likelihood Approximation	Gauss-Hermite Quadrature		
Degrees of Freedom Method	Between-Within		
Fixed Effects SE Adjustment	Sandwich - MBN(df,r=1,d=2)		

	Class Level Information			
Class	Levels	Values		
TXHOSP	35	GA002 GA003 GA004 GA006 GA008 GA009 GA010 GA011 GA013 GA015 GA016 GA018 GA020 GA021 GA024 GA026 GA027 GA029 GA030 GA032 GA034 GA040 GA046 GA048 GA050 GA056 GA059 GA065 GA066 GA069 GA070 GA071 GA308 GAMDO OSODC		
mrsafinal	2	MRSA MSSA		
kidney	2	Yes No		
DIABETES	2	Yes No		
SMOKER	2	Yes No		
WOUND	2	Yes No		
BSI	2	Yes No		

Number of Observations Read	1447
Number of Observations Used	1447

Response Profile			
Ordered Value	Total Frequency		
1	No	735	
2	Yes	712	

The GLIMMIX procedure is modeling the probability that hosp='Yes'.

The GLIMMIX Procedure

Dimensions	
G-side Cov. Parameters	1
Columns in X	14
Columns in Z per Subject	1
Subjects (Blocks in V)	35
Max Obs per Subject	447

Optimization Information			
Optimization Technique Dual Quasi-Newton			
Parameters in Optimization	9		
Lower Boundaries	1		
Upper Boundaries	0		
Fixed Effects	Not Profiled		
Starting From	GLM estimates		
Quadrature Points	1		

Iteration History					
Iteration	Restarts	Evaluations	Objective Function	Change	Max Gradient
0	0	4	2809.7671492		2776.37
1	0	6	2798.1628166	11.60433259	120.7446
2	0	4	2738.8833011	59.27951551	40.08431
3	0	2	2732.1785248	6.70477633	19.49736
4	0	4	2728.8319698	3.34655496	14.21317
5	0	2	2728.0560481	0.77592168	4.021616
6	0	3	2727.5457195	0.51032860	34.32043
7	0	2	2727.0471694	0.49855013	35.08955
8	0	2	2726.3982854	0.64888401	8.700023
9	0	3	2726.1195295	0.27875594	9.737444
10	0	3	2726.0684443	0.05108520	3.665075
11	0	3	2726.0633297	0.00511460	1.240708
12	0	3	2726.0609073	0.00242232	0.580288
13	0	3	2726.0600159	0.00089147	0.146745
14	0	3	2726.0600102	0.00000571	0.011663

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

The GLIMMIX Procedure

Fit Statistics		
-2 Log Likelihood	2726.06	
AIC (smaller is better)	2744.06	
AICC (smaller is better)	2744.19	
BIC (smaller is better)	2758.06	
CAIC (smaller is better)	2767.06	
HQIC (smaller is better)	2748.89	

Fit Statistics for Conditional Distribution			
-2 log L(hosp r. effects) 2605.19			
Pearson Chi-Square	1867.72		
Pearson Chi-Square / DF	1.29		

Covariance Parameter Estimates					
Cov Parm Subject Estimate Standard Z Value Pr >					Pr > Z
Intercept	TXHOSP	3.3699	2.2234	1.52	0.0648

Solutions for Fixed Effects												
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t	
Intercept							-2.2875	0.5930	34	-3.86	0.0005	
SMOKER				Yes			0.5006	0.3209	18	1.56	0.1362	
SMOKER				No			0					
mrsafinal	MRSA						0.01493	0.2164	24	0.07	0.9456	
mrsafinal	MSSA						0					
kidney		Yes					1.2379	0.3360	22	3.68	0.0013	
kidney		No					0					
DIABETES			Yes				0.7665	0.2814	22	2.72	0.0124	
DIABETES			No				0					
WOUND					Yes		0.4193	0.2014	21	2.08	0.0498	
WOUND					No		0					
BSI						Yes	2.7266	0.3766	24	7.24	<.0001	
BSI						No	0					
newage							0.02178	0.005370	4321	4.06	<.0001	

The GLIMMIX Procedure

Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BS
			Yes			44.242				No		
MRSA						44.242	MSSA					
	Yes					44.242		No				
		Yes				44.242			No			
				Yes		44.242					No	
					Yes	44.242						No
						45.242						

Effects of continuous variables are assessed as one unit offsets from the mean.

The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

The GLIMMIX Procedure

Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	_newage
			Yes			44.242				No		44.242
MRSA						44.242	MSSA					44.242
	Yes					44.242		No				44.242
		Yes				44.242			No			44.242
				Yes		44.242					No	44.242
					Yes	44.242						44.242
						45.242						44.242

Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

The GLIMMIX Procedure

Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	Estimate
			Yes			44.242				No		1.650
MRSA						44.242	MSSA					1.015
	Yes					44.242		No				3.448
		Yes				44.242			No			2.152
				Yes		44.242					No	1.521
					Yes	44.242						15.281
						45.242						1.022

Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

The GLIMMIX Procedure

	Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	DF	95% Confi denc e Limit s
			Yes			44.242				No		18	0.841
MRSA						44.242	MSSA					24	0.649
	Yes					44.242		No				22	1.718
		Yes				44.242			No			22	1.201
				Yes		44.242					No	21	1.000
					Yes	44.242						24	7.024
						45.242						4321	1.011

The GLIMMIX Procedure

	Odds Ratio Estimates											
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	95% Confi dence Limits
			Yes			44.242				No		3.238
MRSA						44.242	MSSA					1.587
	Yes					44.242		No				6.923
		Yes				44.242			No			3.858
				Yes		44.242					No	2.312
					Yes	44.242						33.242
						45.242						1.033

	Type III Tests of Fixed Effects									
Effect	Num DF	Den DF	Chi-Square	F Value	Pr > ChiSq	Pr > F				
SMOKER	1	18	2.43	2.43	0.1188	0.1362				
mrsafinal	1	24	0.00	0.00	0.9450	0.9456				
kidney	1	22	13.57	13.57	0.0002	0.0013				
DIABETES	1	22	7.42	7.42	0.0065	0.0124				
WOUND	1	21	4.33	4.33	0.0374	0.0498				
BSI	1	24	52.43	52.43	<.0001	<.0001				
newage	1	4321	16.45	16.45	<.0001	<.0001				

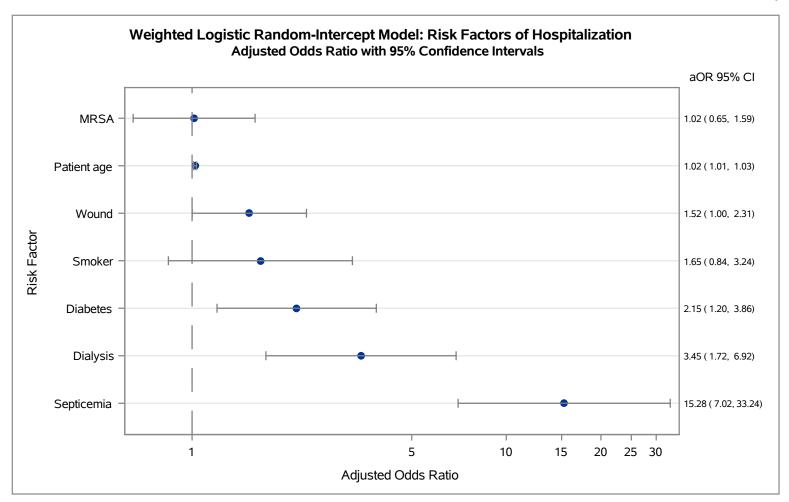
		So	lution for	Randor	n Effects				
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA002	1.0334	0.8120	4355	1.27	0.2032	0.05	-0.5585	2.6253
Intercept	TXHOSP GA003	0.1329	0.6562	4355	0.20	0.8395	0.05	-1.1536	1.4194
Intercept	TXHOSP GA004	1.5172	0.5979	4355	2.54	0.0112	0.05	0.3450	2.6894
Intercept	TXHOSP GA006	0.1466	0.9143	4355	0.16	0.8726	0.05	-1.6460	1.9391
Intercept	TXHOSP GA008	-0.9284	0.6812	4355	-1.36	0.1730	0.05	-2.2638	0.4071
Intercept	TXHOSP GA009	1.6056	0.6720	4355	2.39	0.0169	0.05	0.2882	2.9230
Intercept	TXHOSP GA010	1.1908	0.6076	4355	1.96	0.0501	0.05	-0.00038	2.3820
Intercept	TXHOSP GA011	1.3265	0.5764	4355	2.30	0.0214	0.05	0.1964	2.4566
Intercept	TXHOSP GA013	1.1744	0.7535	4355	1.56	0.1192	0.05	-0.3028	2.6516
Intercept	TXHOSP GA015	1.8955	0.6849	4355	2.77	0.0057	0.05	0.5527	3.2382
Intercept	TXHOSP GA016	0.7896	1.4907	4355	0.53	0.5963	0.05	-2.1328	3.7121

	Solution for Random Effects										
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper		
Intercept	TXHOSP GA018	0.5921	0.6871	4355	0.86	0.3889	0.05	-0.7550	1.9392		
Intercept	TXHOSP GA020	1.0831	0.5938	4355	1.82	0.0682	0.05	-0.08112	2.2473		
Intercept	TXHOSP GA021	1.0662	0.6055	4355	1.76	0.0783	0.05	-0.1208	2.2533		
Intercept	TXHOSP GA024	0.7462	0.5985	4355	1.25	0.2126	0.05	-0.4273	1.9196		
Intercept	TXHOSP GA026	1.3636	0.6238	4355	2.19	0.0289	0.05	0.1407	2.5865		
Intercept	TXHOSP GA027	1.5192	0.6421	4355	2.37	0.0180	0.05	0.2603	2.7781		
Intercept	TXHOSP GA029	-0.3851	1.5919	4355	-0.24	0.8089	0.05	-3.5060	2.7358		
Intercept	TXHOSP GA030	-1.7577	1.1774	4355	-1.49	0.1356	0.05	-4.0660	0.5506		
Intercept	TXHOSP GA032	1.7465	0.6516	4355	2.68	0.0074	0.05	0.4691	3.0239		
Intercept	TXHOSP GA034	-0.4074	0.6659	4355	-0.61	0.5407	0.05	-1.7130	0.8981		
Intercept	TXHOSP GA040	-4.6288	0.8987	4355	-5.15	<.0001	0.05	-6.3908	-2.8668		
Intercept	TXHOSP GA046	0.2668	0.7549	4355	0.35	0.7238	0.05	-1.2133	1.7468		
Intercept	TXHOSP GA048	-0.03924	1.0758	4355	-0.04	0.9709	0.05	-2.1483	2.0698		
Intercept	TXHOSP GA050	-1.7129	1.4468	4355	-1.18	0.2365	0.05	-4.5494	1.1236		
Intercept	TXHOSP GA056	0.3922	1.6040	4355	0.24	0.8068	0.05	-2.7524	3.5368		
Intercept	TXHOSP GA059	0.4726	0.6188	4355	0.76	0.4451	0.05	-0.7406	1.6859		
Intercept	TXHOSP GA065	-3.4060	1.3883	4355	-2.45	0.0142	0.05	-6.1277	-0.6842		
Intercept	TXHOSP GA066	-0.8944	1.3783	4355	-0.65	0.5164	0.05	-3.5965	1.8077		
Intercept	TXHOSP GA069	1.4220	1.4749	4355	0.96	0.3350	0.05	-1.4694	4.3135		
Intercept	TXHOSP GA070	0.8376	0.6511	4355	1.29	0.1984	0.05	-0.4390	2.1141		
Intercept	TXHOSP GA071	-0.1330	0.6855	4355	-0.19	0.8462	0.05	-1.4769	1.2110		
Intercept	TXHOSP GA308	-2.3440	1.3530	4355	-1.73	0.0833	0.05	-4.9966	0.3086		
Intercept	TXHOSP GAMDO	-3.2892	0.5959	4355	-5.52	<.0001	0.05	-4.4574	-2.1210		
Intercept	TXHOSP OSODC	-1.8421	1.4661	4355	-1.26	0.2090	0.05	-4.7164	1.0322		

	Empirical Correlation Matrix for Fixed Effects												
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col1	Col2	Col3	Col4	Col5	Col6
Intercept							1	1.0000	-0.08803		-0.2056		-0.06619
SMOKER				Yes			2	-0.08803	1.0000		0.08159		0.03500
SMOKER				No			3			1.0000			
mrsafinal	MRSA						4	-0.2056	0.08159		1.0000		0.1599
mrsafinal	MSSA						5					1.0000	
kidney		Yes					6	-0.06619	0.03500		0.1599		1.0000
kidney		No					7						
DIABETES			Yes				8	-0.04415	0.2710		0.1505		-0.2076
DIABETES			No				9						
WOUND					Yes		10	-0.1236	0.3643		0.2547		-0.2413
WOUND					No		11						
BSI						Yes	12	-0.07947	0.1116		0.1467		0.02663
BSI						No	13						
newage							14	-0.1829	-0.07769		-0.3066		-0.3054

	Empirical Correlation Matrix for Fixed Effects													
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col7	Col8	Col9	Col10	Col11	Col12	Col13
Intercept							1		-0.04415		-0.1236		-0.07947	
SMOKER				Yes			2		0.2710		0.3643		0.1116	
SMOKER				No			3							
mrsafinal	MRSA						4		0.1505		0.2547		0.1467	
mrsafinal	MSSA						5							
kidney		Yes					6		-0.2076		-0.2413		0.02663	
kidney		No					7	1.0000						
DIABETES			Yes				8		1.0000		0.07682		-0.05413	
DIABETES			No				9			1.0000				
WOUND					Yes		10		0.07682		1.0000		0.2771	
WOUND					No		11					1.0000		
BSI						Yes	12		-0.05413		0.2771		1.0000	
BSI						No	13							1.0000
newage							14		-0.1068		0.03849		-0.04147	

	Empirical Correlation Matrix for Fixed Effects									
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col14		
Intercept							1	-0.1829		
SMOKER				Yes			2	-0.07769		
SMOKER				No			3			
mrsafinal	MRSA						4	-0.3066		
mrsafinal	MSSA						5			
kidney		Yes					6	-0.3054		
kidney		No					7			
DIABETES			Yes				8	-0.1068		
DIABETES			No				9			
WOUND					Yes		10	0.03849		
WOUND					No		11			
BSI						Yes	12	-0.04147		
BSI						No	13			
newage							14	1.0000		



Model Info	ormation
Data Set	WORK.COMM
Response Variable	hosp
Response Distribution	Binary
Link Function	Logit
Variance Function	Default
Variance Matrix Blocked By	TXHOSP
Estimation Technique	Maximum Likelihood
Likelihood Approximation	Gauss-Hermite Quadrature
Degrees of Freedom Method	Between-Within
Fixed Effects SE Adjustment	Sandwich - MBN(df,r=1,d=2)

	Class Level Information								
Class	Levels	Values							
TXHOSP	30	GA002 GA003 GA004 GA006 GA008 GA009 GA010 GA011 GA013 GA015 GA016 GA018 GA020 GA021 GA024 GA026 GA027 GA030 GA032 GA034 GA040 GA046 GA048 GA050 GA059 GA066 GA069 GA070 GA071 GAMDO							
mrsafinal	2	MRSA MSSA							
kidney	2	Yes No							
DIABETES	2	Yes No							
SMOKER	2	Yes No							
WOUND	2	Yes No							
BSI	2	Yes No							

Number of Observations Read	846
Number of Observations Used	846

Response Profile							
Ordered Value	hosp	Total Frequency					
1	No	590					
2	Yes	256					

The GLIMMIX procedure is modeling the probability that hosp='Yes'.

Dimensions	
G-side Cov. Parameters	1
Columns in X	14
Columns in Z per Subject	1
Subjects (Blocks in V)	30
Max Obs per Subject	380

Optimization Info	rmation
Optimization Technique	Dual Quasi-Newton
Parameters in Optimization	9
Lower Boundaries	1
Upper Boundaries	0
Fixed Effects	Not Profiled
Starting From	GLM estimates
Quadrature Points	1

		Iterat	ion History		
Iteration	Restarts	Evaluations	Objective Function	Change	Max Gradient
0	0	4	1501.0917107		2753.628
1	0	6	1480.828851	20.26285972	79.96838
2	0	2	1446.3052458	34.52360519	25.9747
3	0	2	1438.6102748	7.69497099	25.24847
4	0	4	1434.7206514	3.88962333	47.57425
5	0	4	1431.4966568	3.22399465	12.60808
6	0	3	1431.0656799	0.43097691	13.71226
7	0	4	1430.0803129	0.98536704	5.962923
8	0	3	1429.733555	0.34675782	14.80668
9	0	3	1429.6672263	0.06632873	2.606062
10	0	3	1429.6524119	0.01481443	1.815746
11	0	2	1429.6338936	0.01851827	2.527292
12	0	3	1429.6301493	0.00374429	0.276731
13	0	3	1429.6300976	0.00005169	0.184168
14	0	3	1429.6300848	0.00001285	0.016909

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

Fit Statistics	
-2 Log Likelihood	1429.63
AIC (smaller is better)	1447.63
AICC (smaller is better)	1447.85
BIC (smaller is better)	1460.24
CAIC (smaller is better)	1469.24
HQIC (smaller is better)	1451.66

Fit Statistics for Condi Distribution	tional
-2 log L(hosp r. effects)	1329.79
Pearson Chi-Square	1214.52
Pearson Chi-Square / DF	1.44

	Covaria	nce Parame	eter Estimate	es	
Cov Parm	Subject	Estimate	Standard Error	Z Value	Pr > Z
Intercept	TXHOSP	4.1805	2.3097	1.81	0.0351

	Solutions for Fixed Effects													
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t			
Intercept							-2.5290	0.6961	29	-3.63	0.0011			
SMOKER				Yes			0.6718	0.4220	17	1.59	0.1298			
SMOKER				No			0							
mrsafinal	MRSA						-0.1810	0.2851	22	-0.64	0.5320			
mrsafinal	MSSA						0							
kidney		Yes					1.6130	0.6778	10	2.38	0.0386			
kidney		No					0							
DIABETES			Yes				1.1581	0.4643	17	2.49	0.0232			
DIABETES			No				0							
WOUND					Yes		0.5562	0.3572	16	1.56	0.1391			
WOUND					No		0							
BSI						Yes	4.0945	0.7998	19	5.12	<.0001			
BSI						No	0							
newage							0.02049	0.007379	2870	2.78	0.0055			

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BS			
			Yes			39.545				No					
MRSA						39.545	MSSA								
	Yes					39.545		No							
		Yes				39.545			No						
				Yes		39.545					No				
					Yes	39.545						No			
						40.545									

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	_newage			
			Yes			39.545				No		39.545			
MRSA						39.545	MSSA					39.545			
	Yes					39.545		No				39.545			
		Yes				39.545			No			39.545			
				Yes		39.545					No	39.545			
					Yes	39.545						39.545			
						40.545						39.545			

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	Estimate			
			Yes			39.545				No		1.958			
MRSA						39.545	MSSA					0.834			
	Yes					39.545		No				5.018			
		Yes				39.545			No			3.184			
				Yes		39.545					No	1.744			
					Yes	39.545						60.007			
						40.545						1.021			

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	DF	95% Confi dence Limits		
			Yes			39.545				No		17	0.804		
MRSA						39.545	MSSA					22	0.462		
	Yes					39.545		No				10	1.108		
		Yes				39.545			No			17	1.196		
				Yes		39.545					No	16	0.818		
					Yes	39.545						19	11.252		
						40.545						2870	1.006		

					(Odds Ratio	Estimates					
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	95% Confid ence Limits
			Yes			39.545				No		4.768
MRSA						39.545	MSSA					1.507
	Yes					39.545		No				22.721
		Yes				39.545			No			8.480
				Yes		39.545					No	3.719
					Yes	39.545						320.033
						40.545						1.036

Type III Tests of Fixed Effects						
Effect	Num DF	Den DF	F Value	Pr > F		
SMOKER	1	17	2.53	0.1298		
mrsafinal	1	22	0.40	0.5320		
kidney	1	10	5.66	0.0386		
DIABETES	1	17	6.22	0.0232		
WOUND	1	16	2.42	0.1391		
BSI	1	19	26.21	<.0001		
newage	1	2870	7.71	0.0055		

	Solution for Random Effects								
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA002	-1.3852	1.0465	2899	-1.32	0.1857	0.05	-3.4372	0.6668
Intercept	TXHOSP GA003	-0.6542	0.8484	2899	-0.77	0.4407	0.05	-2.3177	1.0093
Intercept	TXHOSP GA004	0.7617	0.6833	2899	1.11	0.2651	0.05	-0.5781	2.1015
Intercept	TXHOSP GA006	-1.3408	1.4341	2899	-0.93	0.3499	0.05	-4.1527	1.4712
Intercept	TXHOSP GA008	-1.6991	0.9333	2899	-1.82	0.0688	0.05	-3.5291	0.1309
Intercept	TXHOSP GA009	1.3989	0.8269	2899	1.69	0.0908	0.05	-0.2224	3.0202
Intercept	TXHOSP GA010	1.2832	0.6953	2899	1.85	0.0651	0.05	-0.08013	2.6465
Intercept	TXHOSP GA011	1.2810	0.6697	2899	1.91	0.0559	0.05	-0.03207	2.5941
Intercept	TXHOSP GA013	2.4816	0.8876	2899	2.80	0.0052	0.05	0.7413	4.2219
Intercept	TXHOSP GA015	2.3732	0.8225	2899	2.89	0.0039	0.05	0.7605	3.9860
Intercept	TXHOSP GA016	0.5821	1.6877	2899	0.34	0.7302	0.05	-2.7270	3.8912
Intercept	TXHOSP GA018	1.0116	0.8259	2899	1.22	0.2207	0.05	-0.6077	2.6310

Solution for Random Effects									
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA020	0.9734	0.6947	2899	1.40	0.1613	0.05	-0.3887	2.3355
Intercept	TXHOSP GA021	0.6896	0.7048	2899	0.98	0.3279	0.05	-0.6923	2.0715
Intercept	TXHOSP GA024	-0.09621	0.7157	2899	-0.13	0.8931	0.05	-1.4995	1.3071
Intercept	TXHOSP GA026	1.4177	0.7398	2899	1.92	0.0554	0.05	-0.03294	2.8683
Intercept	TXHOSP GA027	1.7194	0.7319	2899	2.35	0.0189	0.05	0.2842	3.1545
Intercept	TXHOSP GA030	-1.7086	1.3302	2899	-1.28	0.1991	0.05	-4.3168	0.8996
Intercept	TXHOSP GA032	1.5642	0.7456	2899	2.10	0.0360	0.05	0.1022	3.0262
Intercept	TXHOSP GA034	-1.4768	0.8780	2899	-1.68	0.0927	0.05	-3.1983	0.2447
Intercept	TXHOSP GA040	-4.0165	1.0574	2899	-3.80	0.0001	0.05	-6.0899	-1.9432
Intercept	TXHOSP GA046	2.8069	1.3843	2899	2.03	0.0427	0.05	0.09259	5.5212
Intercept	TXHOSP GA048	-0.7364	1.5905	2899	-0.46	0.6434	0.05	-3.8551	2.3823
Intercept	TXHOSP GA050	-2.4485	1.6212	2899	-1.51	0.1311	0.05	-5.6272	0.7302
Intercept	TXHOSP GA059	0.8361	0.7259	2899	1.15	0.2495	0.05	-0.5872	2.2595
Intercept	TXHOSP GA066	-0.7882	1.5712	2899	-0.50	0.6160	0.05	-3.8690	2.2927
Intercept	TXHOSP GA069	0.3367	1.8053	2899	0.19	0.8520	0.05	-3.2030	3.8764
Intercept	TXHOSP GA070	1.3729	0.8034	2899	1.71	0.0876	0.05	-0.2024	2.9483
Intercept	TXHOSP GA071	-1.0502	0.8703	2899	-1.21	0.2276	0.05	-2.7566	0.6561
Intercept	TXHOSP GAMDO	-4.2320	0.7083	2899	-5.97	<.0001	0.05	-5.6209	-2.8431

Model Information				
Data Set	WORK.HOP			
Response Variable	hosp			
Response Distribution	Binary			
Link Function	Logit			
Variance Function	Default			
Variance Matrix Blocked By	TXHOSP			
Estimation Technique	Maximum Likelihood			
Likelihood Approximation	Gauss-Hermite Quadrature			
Degrees of Freedom Method	Between-Within			
Fixed Effects SE Adjustment	Sandwich - MBN(df,r=1,d=2)			

Class Level Information					
Class	Levels	Values			
TXHOSP	31	GA002 GA003 GA004 GA006 GA008 GA009 GA010 GA011 GA013 GA015 GA018 GA020 GA021 GA024 GA026 GA027 GA029 GA032 GA034 GA040 GA046 GA048 GA056 GA059 GA065 GA069 GA070 GA071 GA308 GAMDO OSODC			
mrsafinal	2	MRSA MSSA			
kidney	2	Yes No			
DIABETES	2	Yes No			
SMOKER	2	Yes No			
WOUND	2	Yes No			
BSI	2	Yes No			

Number of Observations Read	601
Number of Observations Used	601

Response Profile					
Ordered Value	Total Frequency				
1	No	145			
2	Yes	456			

The GLIMMIX procedure is modeling the probability that hosp='Yes'.

Dimensions				
G-side Cov. Parameters	1			
Columns in X	14			
Columns in Z per Subject	1			
Subjects (Blocks in V)	31			
Max Obs per Subject	120			

Optimization Information				
Optimization Technique	Dual Quasi-Newton			
Parameters in Optimization	9			
Lower Boundaries	1			
Upper Boundaries	0			
Fixed Effects	Not Profiled			
Starting From	GLM estimates			
Quadrature Points	5			

Iteration History						
Iteration	Restarts	Evaluations	Objective Function	Change	Max Gradient	
0	0	4	1146.8534811		326.408	
1	0	5	1146.5142917	0.33918935	22.92349	
2	0	4	1135.1806641	11.33362762	8.293453	
3	0	2	1132.1465315	3.03413262	9.172207	
4	0	4	1130.8451183	1.30141320	15.36075	
5	0	4	1129.7493425	1.09577577	2.771987	
6	0	2	1129.3311559	0.41818661	59.56893	
7	0	2	1128.638522	0.69263391	3.230332	
8	0	3	1128.3265204	0.31200159	26.83417	
9	0	2	1128.1767489	0.14977150	41.32122	
10	0	2	1128.127342	0.04940697	49.8322	
11	0	2	1128.0514506	0.07589132	19.01861	
12	0	3	1128.0280862	0.02336444	5.386648	
13	0	3	1128.0140478	0.01403844	0.498978	
14	0	3	1128.0139151	0.00013265	0.16206	
15	0	3	1128.0139063	0.00000880	0.149955	

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

Fit Statistics				
-2 Log Likelihood	1128.01			
AIC (smaller is better)	1146.01			
AICC (smaller is better)	1146.32			
BIC (smaller is better)	1158.92			
CAIC (smaller is better)	1167.92			
HQIC (smaller is better)	1150.22			

Fit Statistics for Conditional Distribution				
-2 log L(hosp r. effects) 1034.77				
Pearson Chi-Square	596.07			
Pearson Chi-Square / DF	0.99			

	Covariance Parameter Estimates										
Cov Parm Subject Estimate Standard Z Value Pr > 2											
Intercept	TXHOSP	4.0459	3.0019	1.35	0.0889						

	Solutions for Fixed Effects											
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t	
Intercept							-0.4119	0.7588	30	-0.54	0.5912	
SMOKER				Yes			0.4916	0.7261	16	0.68	0.5081	
SMOKER				No			0					
mrsafinal	MRSA						0.2375	0.4195	22	0.57	0.5770	
mrsafinal	MSSA						0					
kidney		Yes					0.8999	0.4452	19	2.02	0.0576	
kidney		No					0					
DIABETES			Yes				0.6596	0.3741	19	1.76	0.0940	
DIABETES			No				0					
WOUND					Yes		-0.1607	0.3984	19	-0.40	0.6912	
WOUND					No		0					
BSI						Yes	1.6658	0.5673	21	2.94	0.0079	
BSI						No	0					
newage							0.005249	0.007870	1418	0.67	0.5049	

	Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BSI	
			Yes			53.62				No			
MRSA						53.62	MSSA						
	Yes					53.62		No					
		Yes				53.62			No				
				Yes		53.62					No		
					Yes	53.62						No	
						54.62							

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_newage	
			Yes			53.62				No		53.62	
MRSA						53.62	MSSA					53.62	
	Yes					53.62		No				53.62	
		Yes				53.62			No			53.62	
				Yes		53.62					No	53.62	
					Yes	53.62						53.62	
						54.62						53.62	

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	Estimate	
			Yes			53.62				No		1.635	
MRSA						53.62	MSSA					1.268	
	Yes					53.62		No				2.459	
		Yes				53.62			No			1.934	
				Yes		53.62					No	0.852	
					Yes	53.62						5.290	
						54.62						1.005	

Effects of continuous variables are assessed as one unit offsets from the mean.

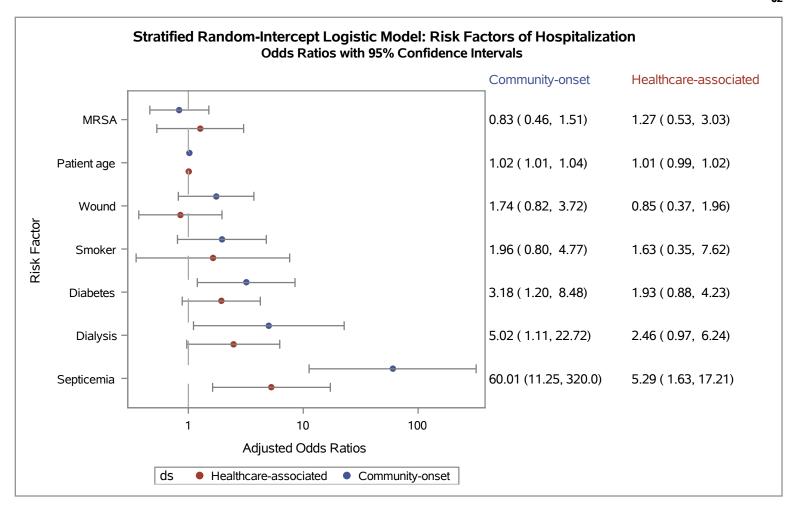
	Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	DF	95% Confi denc e Limit s
			Yes			53.62				No		16	0.351
MRSA						53.62	MSSA					22	0.531
	Yes					53.62		No				19	0.969
		Yes				53.62			No			19	0.884
				Yes		53.62					No	19	0.370
					Yes	53.62						21	1.626
						54.62						1418	0.990

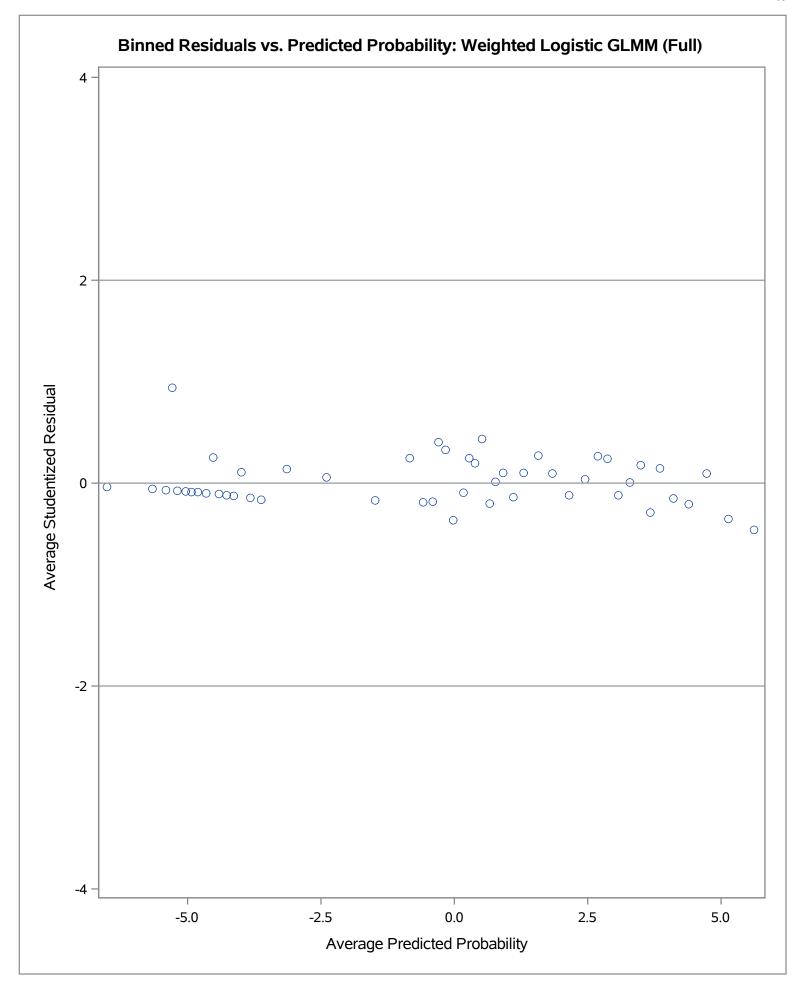
	Odds Ratio Estimates											
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	95% Confi dence Limits
			Yes			53.62				No		7.620
MRSA						53.62	MSSA					3.027
	Yes					53.62		No				6.245
		Yes				53.62			No			4.232
				Yes		53.62					No	1.960
					Yes	53.62						17.212
						54.62						1.021

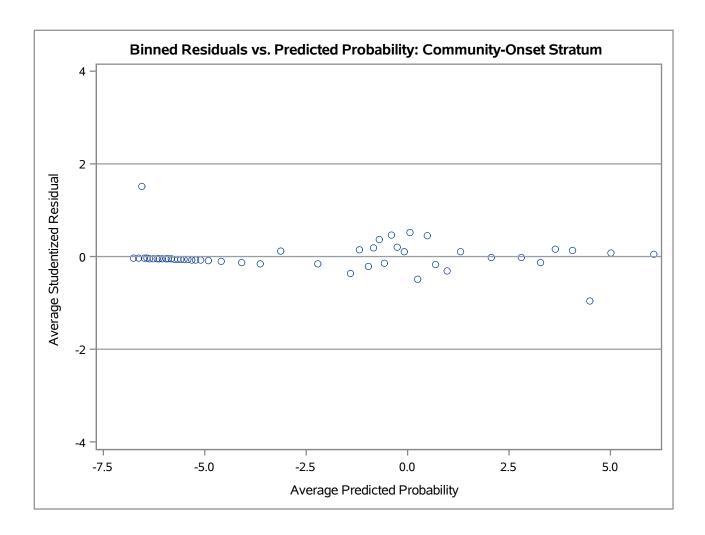
Тур	Type III Tests of Fixed Effects									
Effect	Num DF	Den DF	F Value	Pr > F						
SMOKER	1	16	0.46	0.5081						
mrsafinal	1	22	0.32	0.5770						
kidney	1	19	4.09	0.0576						
DIABETES	1	19	3.11	0.0940						
WOUND	1	19	0.16	0.6912						
BSI	1	21	8.62	0.0079						
newage	1	1418	0.44	0.5049						

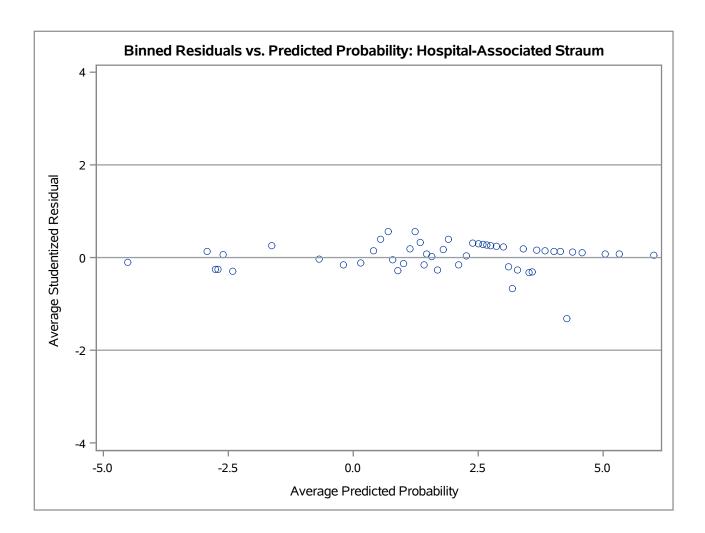
	Solution for Random Effects										
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper		
Intercept	TXHOSP GA002	2.6027	1.3808	1448	1.88	0.0596	0.05	-0.1059	5.3113		
Intercept	TXHOSP GA003	0.5977	0.8245	1448	0.72	0.4686	0.05	-1.0196	2.2150		
Intercept	TXHOSP GA004	1.8489	0.7062	1448	2.62	0.0089	0.05	0.4637	3.2342		
Intercept	TXHOSP GA006	1.3323	1.5733	1448	0.85	0.3972	0.05	-1.7539	4.4186		
Intercept	TXHOSP GA008	-0.9807	0.8354	1448	-1.17	0.2406	0.05	-2.6195	0.6580		
Intercept	TXHOSP GA009	1.0063	0.7729	1448	1.30	0.1932	0.05	-0.5099	2.5224		
Intercept	TXHOSP GA010	0.8390	0.7660	1448	1.10	0.2736	0.05	-0.6636	2.3417		
Intercept	TXHOSP GA011	0.8599	0.6431	1448	1.34	0.1814	0.05	-0.4016	2.1213		
Intercept	TXHOSP GA013	-2.0640	1.1511	1448	-1.79	0.0732	0.05	-4.3219	0.1940		
Intercept	TXHOSP GA015	-0.1339	0.8388	1448	-0.16	0.8732	0.05	-1.7793	1.5115		
Intercept	TXHOSP GA018	0.1910	0.8579	1448	0.22	0.8238	0.05	-1.4918	1.8738		
Intercept	TXHOSP GA020	1.6896	0.7831	1448	2.16	0.0311	0.05	0.1534	3.2258		

	Solution for Random Effects										
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper		
Intercept	TXHOSP GA021	1.0536	0.6838	1448	1.54	0.1236	0.05	-0.2877	2.3949		
Intercept	TXHOSP GA024	0.5776	0.6655	1448	0.87	0.3856	0.05	-0.7279	1.8831		
Intercept	TXHOSP GA026	0.9917	0.7209	1448	1.38	0.1691	0.05	-0.4224	2.4058		
Intercept	TXHOSP GA027	0.02234	0.8926	1448	0.03	0.9800	0.05	-1.7286	1.7733		
Intercept	TXHOSP GA029	-1.0374	1.5837	1448	-0.66	0.5126	0.05	-4.1440	2.0692		
Intercept	TXHOSP GA032	2.9202	1.3270	1448	2.20	0.0279	0.05	0.3170	5.5233		
Intercept	TXHOSP GA034	-0.3605	0.7015	1448	-0.51	0.6074	0.05	-1.7366	1.0156		
Intercept	TXHOSP GA040	-4.5150	1.1279	1448	-4.00	<.0001	0.05	-6.7275	-2.3025		
Intercept	TXHOSP GA046	-1.2222	0.8766	1448	-1.39	0.1634	0.05	-2.9417	0.4972		
Intercept	TXHOSP GA048	0.7424	1.6589	1448	0.45	0.6546	0.05	-2.5117	3.9965		
Intercept	TXHOSP GA056	0.5880	1.6793	1448	0.35	0.7263	0.05	-2.7061	3.8821		
Intercept	TXHOSP GA059	-0.07575	0.7352	1448	-0.10	0.9180	0.05	-1.5179	1.3664		
Intercept	TXHOSP GA065	-3.3498	1.5351	1448	-2.18	0.0293	0.05	-6.3611	-0.3385		
Intercept	TXHOSP GA069	1.0688	1.6253	1448	0.66	0.5109	0.05	-2.1194	4.2571		
Intercept	TXHOSP GA070	0.1242	0.7144	1448	0.17	0.8620	0.05	-1.2772	1.5255		
Intercept	TXHOSP GA071	0.3750	0.9252	1448	0.41	0.6853	0.05	-1.4400	2.1900		
Intercept	TXHOSP GA308	-2.6334	1.4665	1448	-1.80	0.0727	0.05	-5.5100	0.2432		
Intercept	TXHOSP GAMDO	-2.4710	0.6609	1448	-3.74	0.0002	0.05	-3.7674	-1.1746		
Intercept	TXHOSP OSODC	-1.8292	1.5960	1448	-1.15	0.2519	0.05	-4.9600	1.3016		

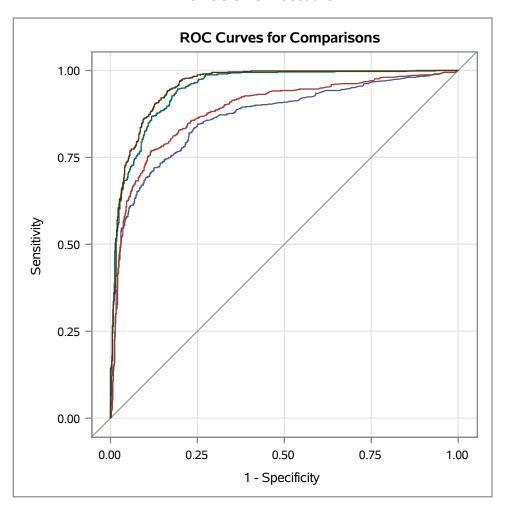








The LOGISTIC Procedure



ROC Association Statistics										
		Mann-	Whitney							
ROC Model	Area	Standard Error	Somers' D	Gamma	Tau-a					
Baseline_Weighted_Logistic_Model	0.8654	0.00980	0.8462	0.8846	0.7309	0.7314	0.3656			
Stratified Weighted Logistic Model	0.8869	0.00900	0.8693	0.9046	0.7739	0.7742	0.3871			
Random-effects Weighted Logistic Model	0.9471	0.00553	0.9362	0.9579	0.8942	0.8942	0.4473			
Stratified, Random-effects Weighted Logistic Model	0.9557	0.00502	0.9459	0.9656	0.9115	0.9115	0.4559			

tion
WORK.TRAINING
HOSPITAL
2
WEIGHT
binary logit
Fisher's scoring

Number of Observations Read	1155
Number of Observations Used	1155
Sum of Weights Read	3510
Sum of Weights Used	3510

Response Profile						
Ordered Value	HOSPITAL	Total Frequency	Total Weight			
1	0	567	2169.0000			
2	1	588	1341.0000			

Probability modeled is HOSPITAL='1'.

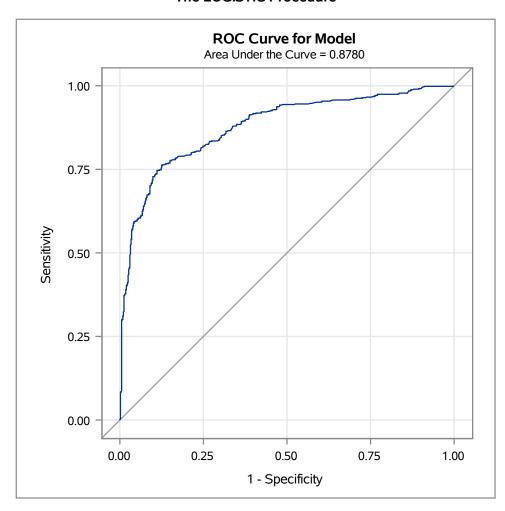
Model Convergence Status
Convergence criterion (GCONV=1E-8) satisfied.

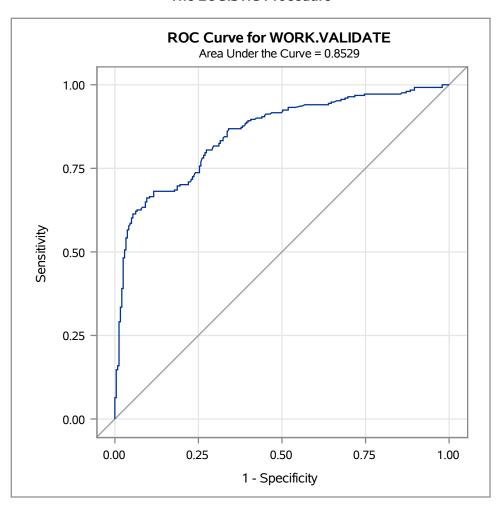
Model Fit Statistics					
Criterion	Intercept and Covariates				
AIC	4670.717	3330.289			
sc	4675.769	3375.756			
-2 Log L	4668.717	3312.289			

Testing Global Null Hypothesis: BETA=0						
Test Chi-Square DF Pr > ChiSq						
Likelihood Ratio	1356.4278	8	<.0001			
Score	1147.9990	8	<.0001			
Wald	685.2433	8	<.0001			

Analysis of Maximum Likelihood Estimates							
Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq		
Intercept	1	-1.1101	0.1331	69.5132	<.0001		
UND	1	-1.0933	0.1128	93.9653	<.0001		
DISLTC8	1	2.0516	0.2517	66.4630	<.0001		
DIABETES	1	0.8131	0.1074	57.3384	<.0001		
RENAL	1	1.2007	0.1559	59.3029	<.0001		
SST	1	-0.7378	0.0919	64.4613	<.0001		
mrsafinal	1	0.8302	0.0919	81.6576	<.0001		
newage	1	0.00818	0.00212	14.8212	0.0001		
BSI	1	2.7088	0.2573	110.8282	<.0001		

Odds Ratio Estimates					
Effect	Point Estimate	95% Wald Confidence Limits			
UND	0.335	0.269	0.418		
DISLTC8	7.780	4.751	12.741		
DIABETES	2.255	1.827	2.783		
RENAL	3.322	2.448 4.510			
SST	0.478	0.399	0.573		
mrsafinal	2.294	1.916	2.746		
newage	1.008	1.004	1.012		
BSI	15.012	9.066	24.857		





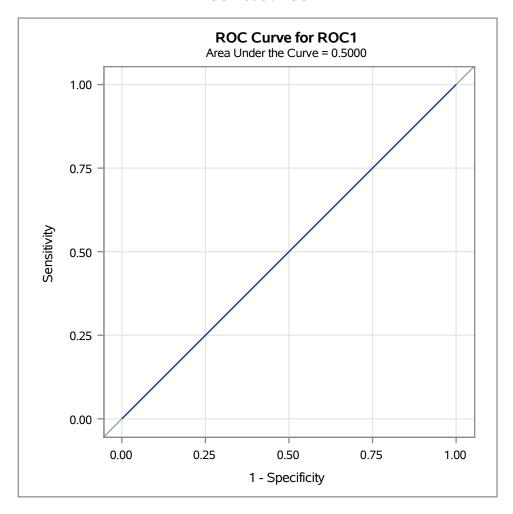
ROC Model: ROC1

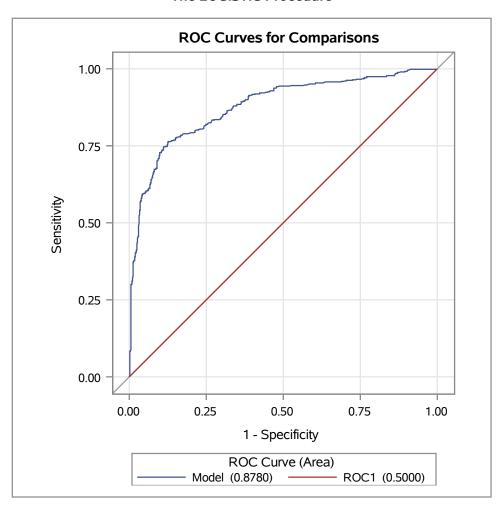
Model Convergence Status Convergence criterion (GCONV=1E-8) satisfied.

> 4668.717 -2 Log L

Analysis of Maximum Likelihood Estimates						
Parameter DF Estimate Standard Wald Chi-Square Pr > ChiSq						
Intercept	1	-0.4809	0.0347	191.6027	<.0001	

ROC Model: ROC1





ROC Association Statistics								
Mann-Whitney								
ROC Model	Area	Standard Error	95% Wald Confidence Limits		Somers' D	Gamma	Tau-a	
Model	0.8780	0.0102	0.8581	0.8980	0.7561	0.7563	0.3782	
	1							

ROC Contrast Test Results					
Contrast DF Chi-Square Pr > ChiSq					
Reference = Model	1	1378.7261	<.0001		

Model Information					
Data Set	WORK.TRAINING				
Response Variable	HOSPITAL				
Number of Response Levels	2				
Weight Variable	WEIGHT				
Model	binary logit				
Optimization Technique	Fisher's scoring				

Number of Observations Read	1155
Number of Observations Used	1155
Sum of Weights Read	3510
Sum of Weights Used	3510

Response Profile				
Ordered Value	HOSPITAL	Total Frequency	Total Weight	
1	0	567	2169.0000	
2	1	588	1341.0000	

Probability modeled is HOSPITAL='1'.

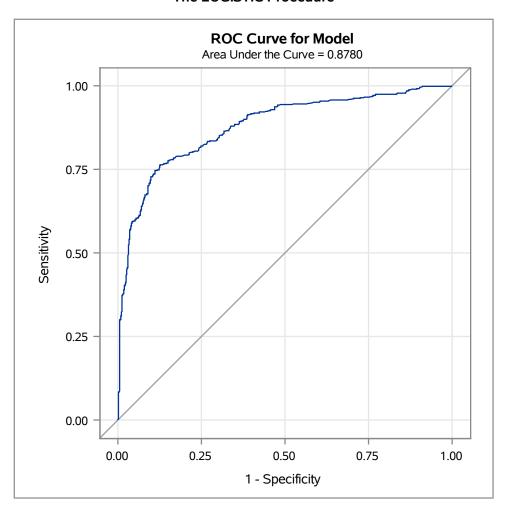
Model Convergence Status
Convergence criterion (GCONV=1E-8) satisfied.

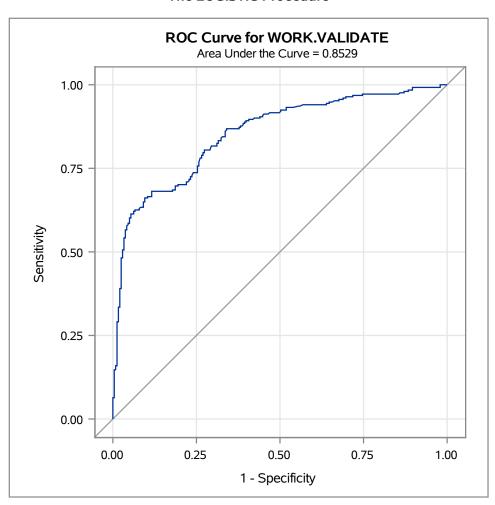
Model Fit Statistics					
Criterion	Intercept and Covariates				
AIC	4670.717	3330.289			
sc	4675.769	3375.756			
-2 Log L	4668.717	3312.289			

Testing Global Null Hypothesis: BETA=0					
Test Chi-Square DF Pr > ChiSq					
Likelihood Ratio	1356.4278	8	<.0001		
Score	1147.9990	8	<.0001		
Wald	685.2433	8	<.0001		

Analysis of Maximum Likelihood Estimates						
Parameter	DF	Estimate	Estimate Standard Chi-Squ		Pr > ChiSq	
Intercept	1	-1.1101	0.1331	69.5132	<.0001	
UND	1	-1.0933	0.1128	93.9653	<.0001	
DISLTC8	1	2.0516	0.2517	66.4630	<.0001	
DIABETES	1	0.8131	0.1074	57.3384	<.0001	
RENAL	1	1.2007	0.1559	59.3029	<.0001	
SST	1	-0.7378	0.0919	64.4613	<.0001	
mrsafinal	1	0.8302	0.0919	81.6576	<.0001	
newage	1	0.00818	0.00212	14.8212	0.0001	
BSI	1	2.7088	0.2573	110.8282	<.0001	

Odds Ratio Estimates				
Effect	Point Estimate	95% Wald Confidence Limits		
UND	0.335	0.269	0.418	
DISLTC8	7.780	4.751	12.741	
DIABETES	2.255	1.827 2.78		
RENAL	3.322	2.448 4.51		
SST	0.478	0.399 0.57		
mrsafinal	2.294	1.916	2.746	
newage	1.008	1.004	1.012	
BSI	15.012	9.066	24.857	





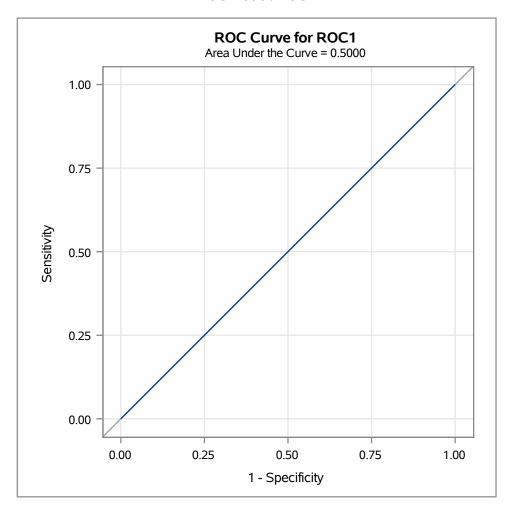
ROC Model: ROC1

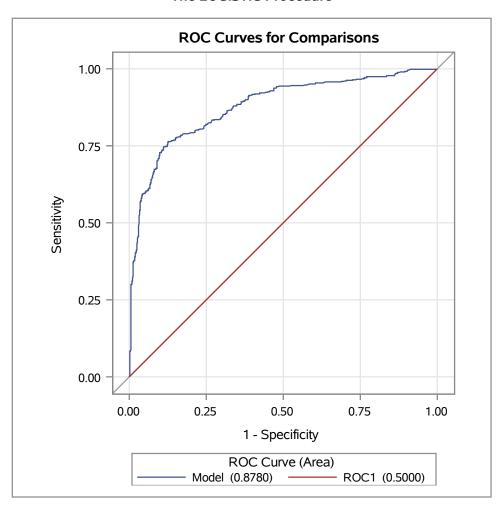
Model Convergence Status Convergence criterion (GCONV=1E-8) satisfied.

> -2 Log L 4668.717

Analysis of Maximum Likelihood Estimates						
Parameter	eter DF Estimate Standard Wald Chi-Square Pr > ChiSq					
Intercept	1	-0.4809	0.0347	191.6027	<.0001	

ROC Model: ROC1





ROC Association Statistics								
Mann-Whitney								
ROC Model	Area	Standard Error	95% Wald Confidence Limits		Somers' D	Gamma	Tau-a	
Model	0.8780	0.0102	0.8581	0.8980	0.7561	0.7563	0.3782	
ROC1	0.5000	0	0.5000	0.5000	0		0	

ROC Contrast Test Results				
Contrast DF Chi-Square Pr > ChiSq				
Reference = Model	1	1378.7261	<.0001	

Model Information				
WORK.TRAINING				
HOSPITAL				
2				
WEIGHT				
binary logit				
Fisher's scoring				

Number of Observations Read	1155
Number of Observations Used	1155
Sum of Weights Read	3510
Sum of Weights Used	3510

Response Profile				
Ordered Value	HOSPITAL	Total Frequency	Total Weight	
1	0	567	2169.0000	
2	1	588	1341.0000	

Probability modeled is HOSPITAL='1'.

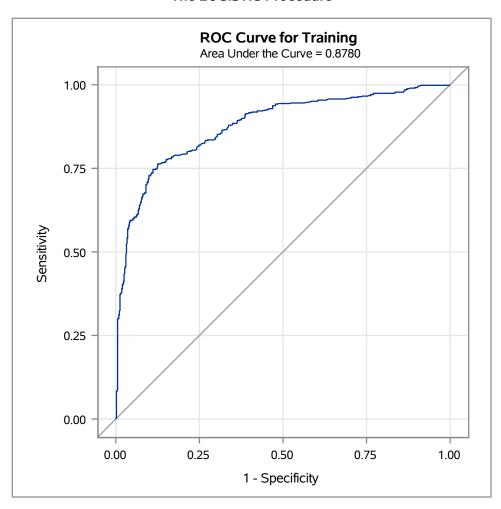
Model Convergence Status
Convergence criterion (GCONV=1E-8) satisfied.

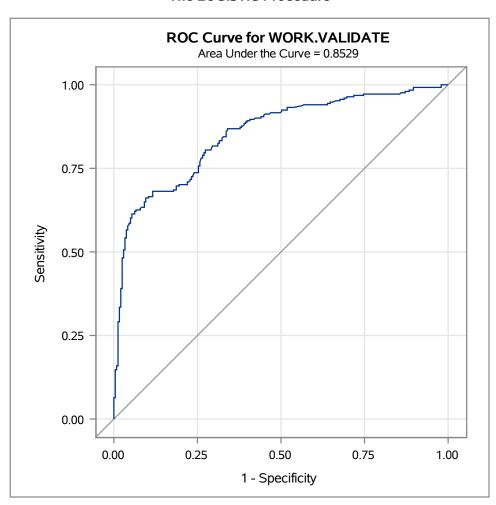
Model Fit Statistics					
Criterion	Intercept Only	Intercept and Covariates			
AIC	4670.717	3330.289			
sc	4675.769	3375.756			
-2 Log L	4668.717	3312.289			

Testing Global Null Hypothesis: BETA=0					
Test	Chi-Square	DF	Pr > ChiSq		
Likelihood Ratio	1356.4278	8	<.0001		
Score	1147.9990	8	<.0001		
Wald	685.2433	8	<.0001		

Analysis of Maximum Likelihood Estimates						
Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq	
Intercept	1	-1.1101	0.1331	69.5132	<.0001	
UND	1	-1.0933	0.1128	93.9653	<.0001	
DISLTC8	1	2.0516	0.2517	66.4630	<.0001	
DIABETES	1	0.8131	0.1074	57.3384	<.0001	
RENAL	1	1.2007	0.1559	59.3029	<.0001	
SST	1	-0.7378	0.0919	64.4613	<.0001	
mrsafinal	1	0.8302	0.0919	81.6576	<.0001	
newage	1	0.00818	0.00212	14.8212	0.0001	
BSI	1	2.7088	0.2573	110.8282	<.0001	

Odds Ratio Estimates				
Effect	Point Estimate	95% Wald Confidence Limits		
UND	0.335	0.269	0.418	
DISLTC8	7.780	4.751	12.741	
DIABETES	2.255	1.827	2.783	
RENAL	3.322	2.448	4.510	
SST	0.478	0.399	0.573	
mrsafinal	2.294	1.916	2.746	
newage	1.008	1.004	1.012	
BSI	15.012	9.066	24.857	





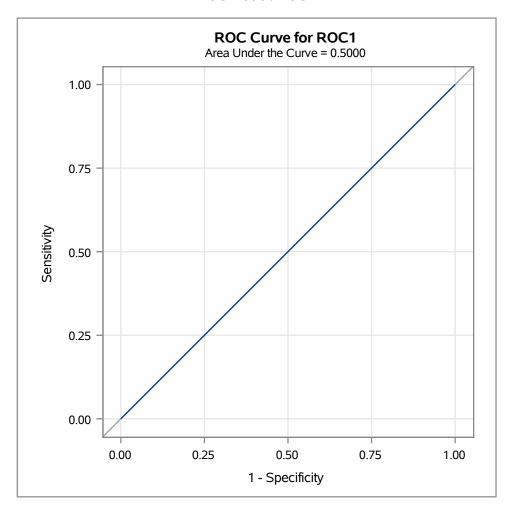
ROC Model: ROC1

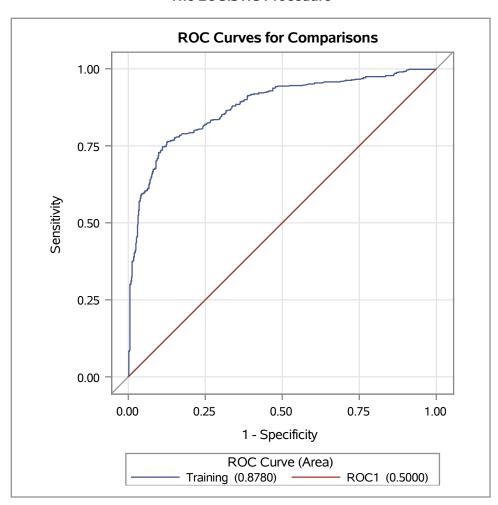
Model Convergence Status Convergence criterion (GCONV=1E-8) satisfied.

> 4668.717 -2 Log L

Analysis of Maximum Likelihood Estimates					
Parameter DF Estimate Standard Wald Chi-Square Pr > ChiSo					
Intercept	1	-0.4809	0.0347	191.6027	<.0001

ROC Model: ROC1





ROC Association Statistics							
	Mann-Whitney						
ROC Model	Area	Standard Error	95% Wald Confidence Limits		Somers' D	Gamma	Tau-a
Training	0.8780	0.0102	0.8581	0.8980	0.7561	0.7563	0.3782
ROC1	0.5000	0	0.5000	0.5000	0		0

ROC Contrast Test Results				
Contrast DF Chi-Square Pr > ChiSq				
Reference = Training	1	1378.7261	<.0001	

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Selecting the Best Model using Honest Assessment

The HPGENSELECT Procedure

Performance Information			
Execution Mode	Single-Machine		
Number of Threads	2		

Data Access Information					
Data Engine Role Path					
WORK.SAMPLE	V9	Input	On Client		
WORK.PRED	V9	Output	On Client		

The HPGENSELECT Procedure

CACASE=0

Model Information				
Data Source	WORK.SAMPLE			
Response Variable	HOSPITAL			
Weight Variable	WEIGHT			
Class Parameterization	GLM			
Distribution	Binary			
Link Function	Logit			
Optimization Technique	Nesterov			

Selection Information				
Selection Method LASSO				
Stop Criterion	None			
Choose Criterion	Validation ASE			
Effect Hierarchy Enforced	None			

Number of Observations						
Description Total Training Validation						
Number of Observations Read	794	557	237			
Number of Observations Used	794	557	237			

Response Profile						
Ordered Value HOSPITAL Frequency Training Validation						
1	0	208	537	238		
2	1	586	917	374		

You are modeling the probability that HOSPITAL='1'.

Class Level Information					
Class	Class Levels Values				
UND	2	1 0			
DISLTC8	2	1 0			
DIABETES	2	1 0			
RENAL	2	1 0			
SST	2	1 0			
mrsafinal	2	1 0			
BSI	2	10			

The HPGENSELECT Procedure

CACASE=0

	Selection Details										
Step	Description	Effects In Model	Lambda	AIC	AICC	BIC	ASE	Validation AIC	Validation AICC	Validation BIC	Validation ASE
0	Initial Model	1	1	1917.197	1917.204	1921.520	0.233	820.935	820.952	824.403	0.238
1	newage entered	2	0.8	1916.144	1916.165	1924.789	0.232	822.841	822.892	829.777	0.238
2		2	0.64	1913.508	1913.529	1922.153	0.232	822.770	822.821	829.706	0.238
3		2	0.512	1911.107	1911.128	1919.752	0.232	822.674	822.726	829.610	0.238
4		2	0.4096	1908.842	1908.864	1917.487	0.231	822.536	822.588	829.472	0.238
5		2	0.3277	1906.665	1906.686	1915.310	0.231	822.355	822.406	829.291	0.238
6		2	0.2621	1904.554	1904.575	1913.199	0.231	822.136	822.187	829.072	0.238
7		2	0.2097	1902.504	1902.526	1911.149	0.230	821.890	821.942	828.826	0.238
8		2	0.1678	1900.517	1900.539	1909.162	0.230	821.628	821.679	828.564	0.237
9		2	0.1342	1898.598	1898.619	1907.243	0.230	821.358	821.410	828.294	0.237
10		2	0.1074	1896.751	1896.773	1905.396	0.229	821.090	821.141	828.026	0.237
11		2	0.0859	1894.982	1895.004	1903.627	0.229	820.829	820.880	827.765	0.237
12		2	0.0687	1893.294	1893.315	1901.939	0.229	820.581	820.632	827.517	0.237
13		2	0.055	1891.688	1891.710	1900.333	0.228	820.350	820.401	827.286	0.237
14		2	0.044	1890.158	1890.180	1898.803	0.228	820.136	820.187	827.072	0.237
15		2	0.0352	1888.712	1888.733	1897.357	0.228	819.943	819.994	826.879	0.237
16	BSI entered	3	0.0281	1888.972	1889.045	1906.262	0.227	822.875	823.047	836.747	0.236
17	UND entered	8	0.0225	1893.776	1894.550	1954.291	0.225	837.822	839.714	886.375	0.234
	DISLTC8 entered	8	0.0225	1893.776	1894.550	1954.291	0.225	837.822	839.714	886.375	0.234
	DIABETES entered	8	0.0225	1893.776	1894.550	1954.291	0.225	837.822	839.714	886.375	0.234
	RENAL entered	8	0.0225	1893.776	1894.550	1954.291	0.225	837.822	839.714	886.375	0.234
	mrsafinal entered	8	0.0225	1893.776	1894.550	1954.291	0.225	837.822	839.714	886.375	0.234
18		8	0.018	1867.880	1868.655	1928.396	0.221	828.627	830.518	877.179	0.231
19		8	0.0144	1837.579	1838.354	1898.095	0.216	817.718	819.610	866.271	0.227
20		8	0.0115	1806.241	1807.016	1866.757	0.211	806.375	808.267	854.928	0.223*

* Optimal Value of Criterion

Maximum Regularization Parameter	4.912624
Chosen Regularization Parameter	0.056639

The HPGENSELECT Procedure

Selected Model

CACASE=0

Selected Effects: Intercept UND DISLTC8 DIABETES RENAL mrsafinal n
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Dimensions			
Number of Effects	8		
Number of Parameters 14			
Columns in X	14		

Fit Statistics				
Training Valida				
-2 Log Likelihood	1778.24	778.38		
AIC (smaller is better)	1806.24	806.38		
AICC (smaller is better)	1807.02	808.27		
BIC (smaller is better)	1866.76	854.93		
Pearson Chi-Square	1353.02	596.07		
Pearson Chi-Square/DF	2.4918	2.6972		
Average Square Error	0.2111	0.2227		

Parameter Estimates					
Parameter	DF	Estimate			
Intercept	1	0.133321			
UND 1	1	-0.053741			
UND 0	1	0.041276			
DISLTC8 1	1	0.032164			
DISLTC8 0	1	-0.042794			
DIABETES 1	1	0.051804			
DIABETES 0	1	-0.068195			
RENAL 1	1	0.060869			
RENAL 0	1	-0.079084			
mrsafinal 1	1	0.057472			
mrsafinal 0	1	-0.074550			
newage	1	0.012053			
BSI 1	1	0.095228			
BSI 0	1	-0.120818			

The HPGENSELECT Procedure

CACASE=1

Model Information				
Data Source	WORK.SAMPLE			
Response Variable	HOSPITAL			
Weight Variable	WEIGHT			
Class Parameterization	GLM			
Distribution	Binary			
Link Function	Logit			
Optimization Technique	Nesterov			

Selection Information				
Selection Method LASSO				
Stop Criterion	None			
Choose Criterion	Validation ASE			
Effect Hierarchy Enforced	None			

Number of Observations						
Description Total Training Valid						
Number of Observations Read	853	598	255			
Number of Observations Used	853	598	255			

Response Profile						
Ordered Value	HOSPITAL	Total Frequency	Training	Validation		
1	0	600	1632	693		
2	1	253	424	186		

You are modeling the probability that HOSPITAL='1'.

Class Level Information					
Class	Levels Value				
UND	2	1 0			
DISLTC8	2	1 0			
DIABETES	2	1 0			
RENAL	2	1 0			
SST	2	1 0			
mrsafinal	2	10			
BSI	2	10			

The HPGENSELECT Procedure

CACASE=1

	Selection Details										
Step	Description	Effects In Model	Lambda	AIC	AICC	BIC	ASE	Validation AIC	Validation AICC	Validation BIC	Validation ASE
0	Initial Model	1	1	2094.650	2094.656	2099.043	0.164	909.413	909.429	912.954	0.167
1	newage entered	2	0.8	2091.491	2091.511	2100.278	0.163	909.182	909.230	916.264	0.166
2		2	0.64	2087.349	2087.369	2096.136	0.163	907.457	907.504	914.539	0.166
3		2	0.512	2083.808	2083.828	2092.595	0.163	906.042	906.090	913.125	0.166
4		2	0.4096	2080.652	2080.672	2089.439	0.162	904.832	904.879	911.914	0.166
5		2	0.3277	2077.770	2077.790	2086.558	0.162	903.767	903.815	910.850	0.165
6		2	0.2621	2075.108	2075.128	2083.895	0.162	902.816	902.864	909.898	0.165
7		2	0.2097	2072.637	2072.658	2081.425	0.162	901.960	902.008	909.043	0.165
8		2	0.1678	2070.345	2070.365	2079.132	0.162	901.189	901.237	908.272	0.165
9		2	0.1342	2068.223	2068.243	2077.010	0.161	900.495	900.542	907.577	0.165
10		2	0.1074	2066.265	2066.285	2075.052	0.161	899.871	899.919	906.954	0.165
11		2	0.0859	2062.857	2062.877	2071.644	0.161	898.834	898.881	905.916	0.165
12		2	0.0687	2060.033	2060.053	2068.820	0.161	898.021	898.068	905.103	0.164
13		2	0.055	2057.697	2057.717	2066.484	0.161	897.390	897.438	904.472	0.164
14		2	0.044	2055.767	2055.787	2064.554	0.160	896.907	896.955	903.989	0.164
15		2	0.0352	2054.174	2054.194	2062.961	0.160	896.543	896.590	903.625	0.164
16	UND entered	4	0.0281	2052.191	2052.333	2078.552	0.160	901.142	901.481	922.390	0.164
	BSI entered	4	0.0281	2052.191	2052.333	2078.552	0.160	901.142	901.481	922.390	0.164
17	DIABETES entered	5	0.0225	2043.048	2043.292	2078.196	0.158	900.297	900.883	928.628	0.163
18	mrsafinal entered	6	0.018	2026.302	2026.676	2070.238	0.157	896.593	897.495	932.006	0.161
19	DISLTC8 entered	7	0.0144	2004.149	2004.683	2056.872	0.154	891.008	892.298	933.504	0.159
20		7	0.0115	1974.349	1974.882	2027.072	0.152	880.283	881.572	922.778	0.157*

* Optimal Value of Criterion

Maximum Regularization Parameter	4.852855
Chosen Regularization Parameter	0.05595

The HPGENSELECT Procedure

Selected Model

CACASE=1

Selected Effects: Intercept UND DISLTC8 DIABETES mrsafinal newage BSI

Dimensions				
Number of Effects 7				
Number of Parameters	12			
Columns in X	12			

Fit Statistics					
Training Valida					
-2 Log Likelihood	1950.35	856.28			
AIC (smaller is better)	1974.35	880.28			
AICC (smaller is better)	1974.88	881.57			
BIC (smaller is better)	2027.07	922.78			
Pearson Chi-Square	1845.09	830.26			
Pearson Chi-Square/DF	3.1486	3.4739			
Average Square Error	0.1519	0.1573			

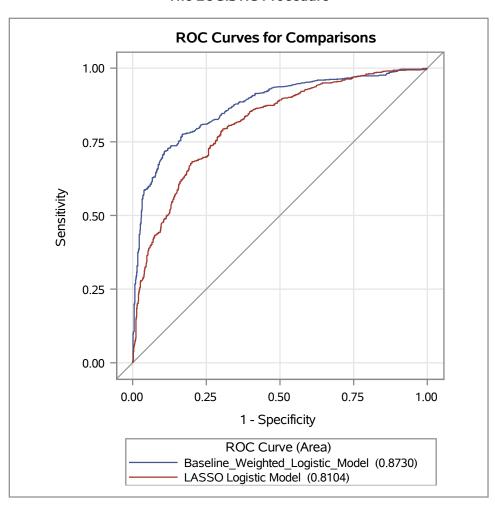
Parameter Estimates					
Parameter	DF	Estimate			
Intercept	1	-1.818762			
UND 1	1	-0.145528			
UND 0	1	0.125829			
DISLTC8 1	1	0.006658			
DISLTC8 0	1	-0.008017			
DIABETES 1	1	0.051218			
DIABETES 0	1	-0.060262			
mrsafinal 1	1	0.024260			
mrsafinal 0	1	-0.028861			
newage	1	0.014599			
BSI 1	1	0.087698			
BSI 0	1	-0.101538			

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Selecting the Best Model using Honest Assessment

The HPGENSELECT Procedure

Procedure Task Timing						
Task Seconds Percent						
Reading and Levelizing Data	0.00	0.43%				
Candidate model fit	0.00	0.57%				
Performing Model Selection	0.34	98.92%				
Producing Output Data Set	0.00	0.08%				



ROC Association Statistics							
		Mann-Whitney					
ROC Model	Area	Standard Error	95% Confiden		Somers' D	Gamma	Tau-a
Baseline_Weighted_Logistic_Model	0.8730	0.00867	0.8560	0.8900	0.7459	0.7462	0.3730
LASSO Logistic Model	0.8104	0.0105	0.7899	0.8310	0.6209	0.6211	0.3105

CACASE=0

Model Information					
Data Set	WORK.PRED				
Response Variable	HOSPITAL				
Number of Response Levels	2				
Weight Variable	WEIGHT				
Model	binary logit				
Optimization Technique	Fisher's scoring				

Number of Observations Read	794
Number of Observations Used	794
Sum of Weights Read	2066
Sum of Weights Used	2066

Response Profile			
Ordered Value	HOSPITAL	Total Frequency	Total Weight
1	0	208	775.0000
2	1	586	1291.0000

Probability modeled is HOSPITAL='1'.

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

Model Fit Statistics				
Intercept and Criterion Only Covariates				
AIC	2735.835	2193.660		
sc	2740.512	2235.753		
-2 Log L	2733.835	2175.660		

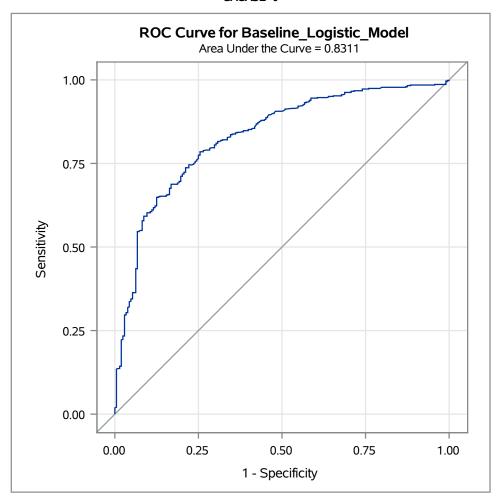
Testing Global Null Hypothesis: BETA=0					
Test Chi-Square DF Pr > ChiSq					
Likelihood Ratio	558.1751	8	<.0001		
Score	452.8053	8	<.0001		
Wald	329.0759	8	<.0001		

CACASE=0

Analysis of Maximum Likelihood Estimates					
Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept	1	0.0450	0.1579	0.0812	0.7757
UND	1	-0.8808	0.1304	45.6112	<.0001
DISLTC8	1	1.8961	0.2667	50.5516	<.0001
DIABETES	1	0.8154	0.1252	42.3977	<.0001
RENAL	1	0.6457	0.1675	14.8667	0.0001
SST	1	-0.3020	0.1104	7.4827	0.0062
mrsafinal	1	0.6834	0.1102	38.4406	<.0001
newage	1	-0.00307	0.00257	1.4254	0.2325
BSI	1	2.1117	0.2645	63.7463	<.0001

Odds Ratio Estimates				
Effect	Point Estimate	95% Wald Confidence Limits		
UND	0.414	0.321	0.535	
DISLTC8	6.660	3.949	11.232	
DIABETES	2.260	1.768	2.889	
RENAL	1.907	1.374	2.648	
SST	0.739	0.595	0.918	
mrsafinal	1.981	1.596	2.458	
newage	0.997	0.992	1.002	
BSI	8.262	4.920	13.874	

CACASE=0

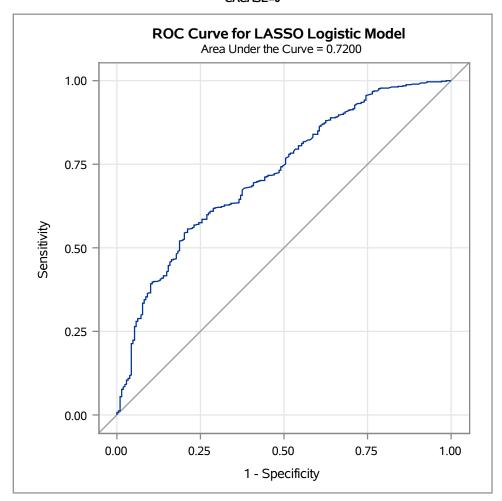


ROC Model: LASSO Logistic Model

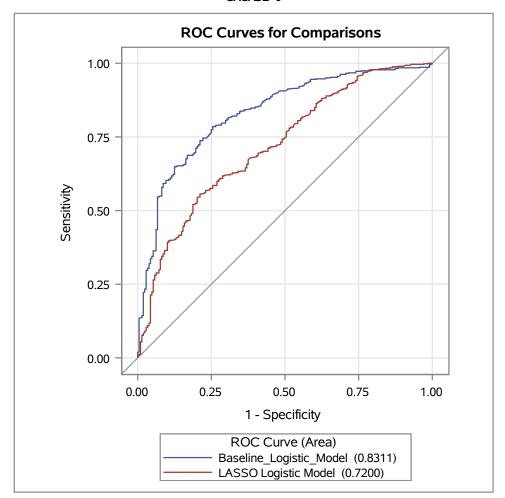
ROC Model Informatio	n
ROC Contrast Coefficients	phat

ROC Model: LASSO Logistic Model

CACASE=0



CACASE=0



ROC Association Statistics							
	Mann-Whitney						
ROC Model	Area	Standard Error			Somers' D	Gamma	Tau-a
Baseline_Logistic_Model	0.8311	0.0161	0.7994	0.8627	0.6622	0.6625	0.2564
LASSO Logistic Model	0.7200	0.0203	0.6802	0.7597	0.4399	0.4403	0.1703

CACASE=1

Model Information			
Data Set WORK.PR			
Response Variable	HOSPITAL		
Number of Response Levels	2		
Weight Variable	WEIGHT		
Model binary logit			
Optimization Technique	Fisher's scoring		

Number of Observations Read	853
Number of Observations Used	853
Sum of Weights Read	2935
Sum of Weights Used	2935

Response Profile				
Ordered Value	HOSPITAL	Total Frequency	Total Weight	
1	0	600	2325.0000	
2	1	253	610.0000	

Probability modeled is HOSPITAL='1'.

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

Model Fit Statistics							
Intercept and Criterion Only Covariate							
AIC	3002.016	2248.671					
sc	3006.765	2291.410					
-2 Log L	3000.016	2230.671					

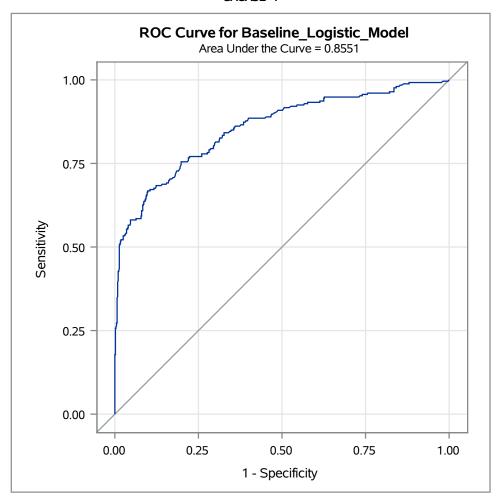
Testing Global Null Hypothesis: BETA=0								
Test Chi-Square DF Pr > ChiSq								
Likelihood Ratio	769.3450	8	<.0001					
Score 820.2740 8 <.00								
Wald	365.7420	8	<.0001					

CACASE=1

Analysis of Maximum Likelihood Estimates									
Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq				
Intercept	1	-1.7720	0.1739	103.8449	<.0001				
UND	1	-1.0551	0.1461	52.1687	<.0001				
DISLTC8	1	3.8047	0.5597	46.2168	<.0001				
DIABETES	1	1.2291	0.1382	79.1148	<.0001				
RENAL	1	0.7382	0.2758	7.1651	0.0074				
SST	1	-0.1288	0.1258	1.0474	0.3061				
mrsafinal	1	0.5465	0.1192	21.0223	<.0001				
newage	1	0.00176	0.00268	0.4329	0.5106				
BSI	1	4.0432	0.3828	111.5694	<.0001				

Odds Ratio Estimates							
Effect	Point Estimate	95% Wald Confidence Limits					
UND	0.348	0.261 0.46					
DISLTC8	44.914	14.997	134.513				
DIABETES	3.418	2.607	4.481				
RENAL	2.092	1.219	3.592				
SST	0.879	0.687	1.125				
mrsafinal	1.727	1.367 2.183					
newage	1.002	0.997 1.007					
BSI	57.006	26.922	120.711				

CACASE=1

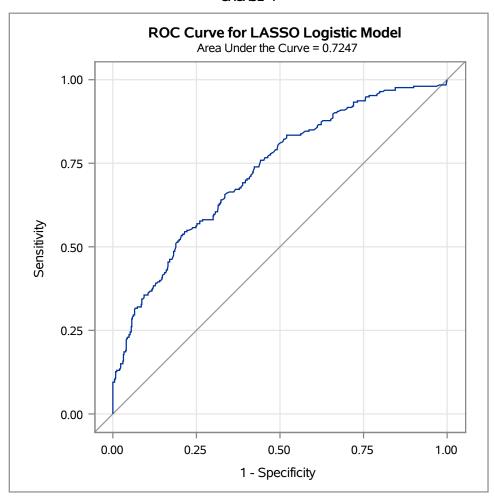


ROC Model: LASSO Logistic Model

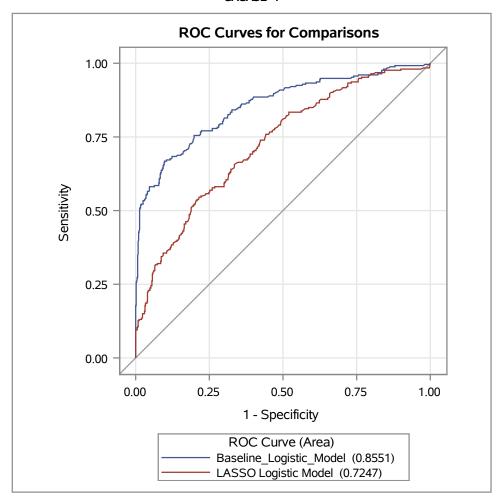
ROC Model Informatio	n
ROC Contrast Coefficients	phat

ROC Model: LASSO Logistic Model

CACASE=1



CACASE=1



ROC Association Statistics								
	Mann-Whitney							
ROC Model	Area	Standard 95% Wald ea Error Confidence Limits			Somers' D	Gamma	Tau-a	
Baseline_Logistic_Model	0.8551	0.0153	0.8251	0.8851	0.7102	0.7106	0.2967	
LASSO Logistic Model	0.7247	0.0190	0.6876	0.7619	0.4495	0.4501	0.1878	

The SURVEYFREQ Procedure

Data Summary	
Number of Observations	1447
Sum of Weights	4363

	Table of hosp by hosp_num										
hosp	hosp_num	Frequency	Weighted Frequency	Std Err of Wgt Freq	Percent	Std Err of Percent					
No	0	735	2808	75.19838	64.3594	1.3469					
	1	0									
	Total	735	2808	75.19838	64.3594	1.3469					
Yes	0	0									
	1	712	1555	57.08056	35.6406	1.3469					
	Total	712	1555	57.08056	35.6406	1.3469					
Total	0	735	2808	75.19838	64.3594	1.3469					
	1	712	1555	57.08056	35.6406	1.3469					
	Total	1447	4363	53.60645	100.0000						

Table of co								
со	Frequency	Weighted Frequency	Std Err of Wgt Freq	Percent	Std Err of Percent			
Community-onset	846	2907	72.90100	66.6285	1.3382			
Healthcare-associated	601	1456	58.42348	33.3715	1.3382			
Total	1447	4363	53.60645	100.0000				

The FREQ Procedure

hosp	hosp_num	Frequency	Percent	Cumulative Frequency	Cumulative Percent
No	0	2808	64.36	2808	64.36
Yes	1	1555	35.64	4363	100.00

со	Frequency	Percent	Cumulative Frequency	Cumulative Percent
Community-onset	2907	66.63	2907	66.63
Healthcare-associated	1456	33.37	4363	100.00

The FREQ Procedure

со	Frequency	Percent	Cumulative Frequency	Cumulative Percent
Community-onset	2907	100.00	2907	100.00

The FREQ Procedure

со	Frequency	Percent	Cumulative Frequency	Cumulative Percent
Healthcare-associated	1456	100.00	1456	100.00

SAS Modeling Output Baseline Weighted Logistic Regression Model

The SURVEYLOGISTIC Procedure

Model Information		
Data Set	WORK.STAPH	
Response Variable	hosp	
Number of Response Levels	2	
Stratum Variable	INVASIVE	
Number of Strata	2	
Weight Variable	WEIGHT	
Model	Binary Logit	
Optimization Technique	Newton-Raphson	
Variance Adjustment	Degrees of Freedom (DF)	

Variance Estimation		
Method	Taylor Series	
Variance Adjustment	Degrees of Freedom (DF)	

Number of Observations Read	1447
Number of Observations Used	1447
Sum of Weights Read	4363
Sum of Weights Used	4363

Response Profile			
Ordered Value	hosp	Total Frequency	Total Weight
1	No	735	2808.0000
2	Yes	712	1555.0000

Probability modeled is hosp='Yes'.

Class Level Information			
Value	Design Variables		
No	0		
Yes	1		
MRSA	1		
MSSA	0		
No	0		
Yes	1		
No	0		
Yes	1		
	Value No Yes MRSA MSSA No Yes No		

SAS Modeling Output Baseline Weighted Logistic Regression Model

The SURVEYLOGISTIC Procedure

Class Level Information				
Class	Value	Design Variables		
SMOKER	No	0		
	Yes	1		
BSI	No	0		
	Yes	1		

Stratum Information				
Stratum INVASIVE N Obs				
1	No	953		
2	Yes	494		

Model Convergence Status

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

Model Fit Statistics						
Criterion	Criterion Intercept Only					
AIC	5685.439	4134.144				
sc	5691.820	4185.191				
-2 Log L	5683.439	4118.144				

	R-Square	0.3015	Max-rescaled R-Square	0.4140	
ı	- 1				ı

Testing Global Null Hypothesis: BETA=0						
Test	F Value	Num DF	Den DF	Pr > F		
Likelihood Ratio	74.16	7.0000	10115	<.0001		
Score	121.72	7	1439	<.0001		
Wald	39.21	7	1439	<.0001		

NOTE:

Second-order Rao-Scott design correction 0.0000 applied to the Likelihood Ratio test.

SAS Modeling Output Baseline Weighted Logistic Regression Model

The SURVEYLOGISTIC Procedure

Type 3 Analysis of Effects							
Effect	Den DF	Pr > F					
SMOKER	22.54	1	1445	<.0001			
mrsafinal	22.41	1	1445	<.0001			
kidney	13.38	1	1445	0.0003			
DIABETES	35.62	1	1445	<.0001			
BSI	158.48	1	1445	<.0001			
WOUND	15.02	1	1445	0.0001			
newage	9.69	1	1445	0.0019			

Analysis of Maximum Likelihood Estimates						
Parameter		Estimate	Standard Error	t Value	Pr > t	
Intercept		-2.3264	0.1864	-12.48	<.0001	
SMOKER	Yes	1.0095	0.2126	4.75	<.0001	
mrsafinal	MRSA	0.7667	0.1620	4.73	<.0001	
kidney	Yes	1.0379	0.2838	3.66	0.0003	
DIABETES	Yes	1.1657	0.1953	5.97	<.0001	
BSI	Yes	3.6303	0.2884	12.59	<.0001	
WOUND	Yes	0.6579	0.1697	3.88	0.0001	
newage		0.0108	0.00346	3.11	0.0019	
NOTE: 1	The degre	ees of freed	om for the t	tests is 1	445.	

Odds Ratio Estimates					
Effect	Point Estimate	95% Confidence Limits			
SMOKER Yes vs No	2.744	1.808	4.164		
mrsafinal MRSA vs MSSA	2.153	1.567	2.958		
kidney Yes vs No	2.823	1.618	4.926		
DIABETES Yes vs No	3.208	2.187	4.706		
BSI Yes vs No	37.725	21.427	66.420		
WOUND Yes vs No	1.931	1.384	2.694		
newage	1.011	1.004	1.018		

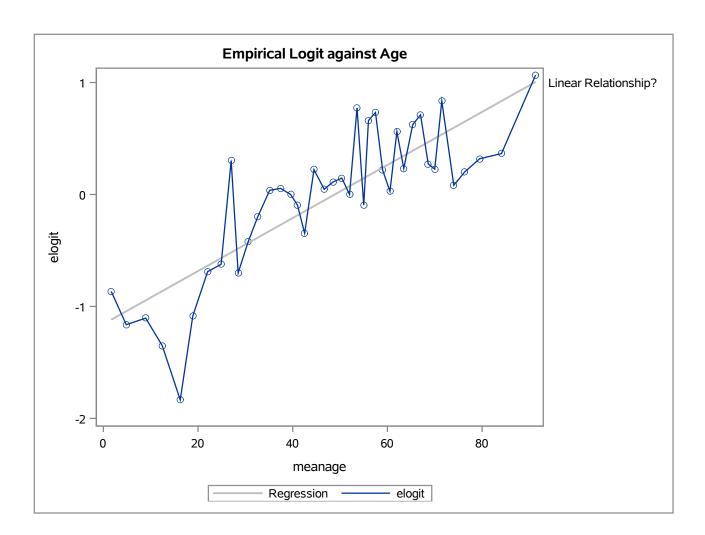
NOTE: The degrees of freedom in computing the confidence limits is 1445.

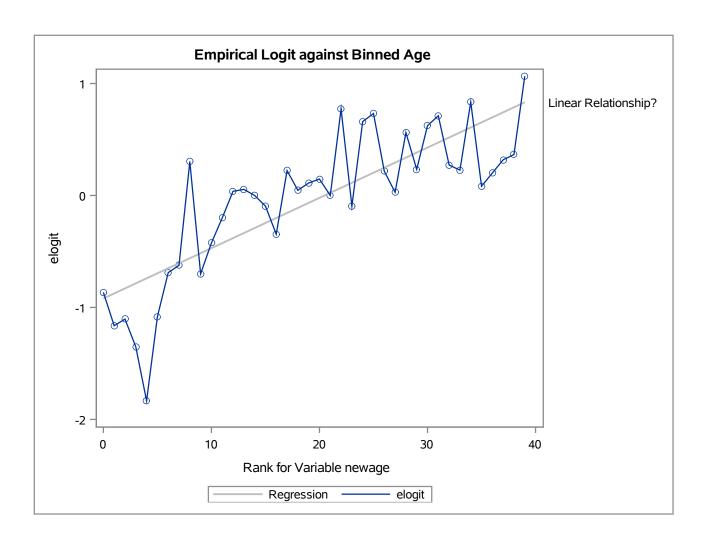
SAS Modeling Output Baseline Weighted Logistic Regression Model

The SURVEYLOGISTIC Procedure

Association of Predicted Probabilities and Observed Responses						
Percent Concordant 86.4 Somers' D 0.731						
Percent Discordant	13.4	Gamma	0.732			
Percent Tied	Tau-a	0.366				
Pairs	523320	С	0.865			

	Estimated Correlation Matrix							
Parameter	Intercept	SMOKERYes	mrsafinalMRSA	kidneyYes	DIABETESYes	BSIYes	WOUNDYes	newage
Intercept	1.0000	-0.1785	-0.3593	0.0276	0.0005	-0.1332	-0.1622	-0.7818
SMOKERYes	-0.1785	1.0000	-0.0766	0.0801	0.0184	0.0226	-0.0322	0.0285
mrsafinalMRSA	-0.3593	-0.0766	1.0000	-0.0583	-0.0496	0.0480	0.0557	0.1120
kidneyYes	0.0276	0.0801	-0.0583	1.0000	-0.1249	-0.0402	0.0667	-0.1625
DIABETESYes	0.0005	0.0184	-0.0496	-0.1249	1.0000	0.1465	-0.0995	-0.1909
BSIYes	-0.1332	0.0226	0.0480	-0.0402	0.1465	1.0000	0.1665	-0.0821
WOUNDYes	-0.1622	-0.0322	0.0557	0.0667	-0.0995	0.1665	1.0000	-0.1344
newage	-0.7818	0.0285	0.1120	-0.1625	-0.1909	-0.0821	-0.1344	1.0000





The CORR Procedure

2 Variables: elogit meanage

Simple Statistics						
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum
elogit	40	-0.04369	0.65292	-1.74763	-1.83438	1.06538
meanage	40	47.02262	22.85461	1881	1.65789	91.29730

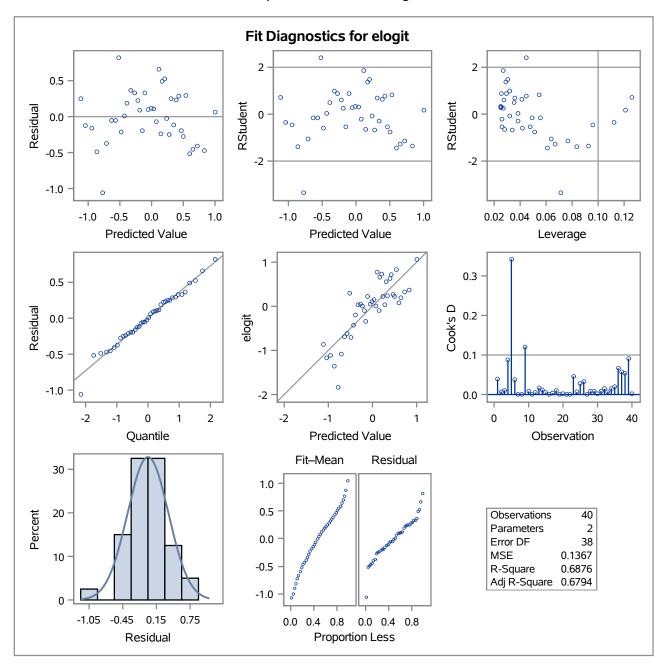
Pearson Correlation Coefficients, N = 40 Prob > r under H0: Rho=0					
	elogit meanage				
elogit	1.00000	0.82924 <.0001			
meanage	0.82924 <.0001	1.00000			

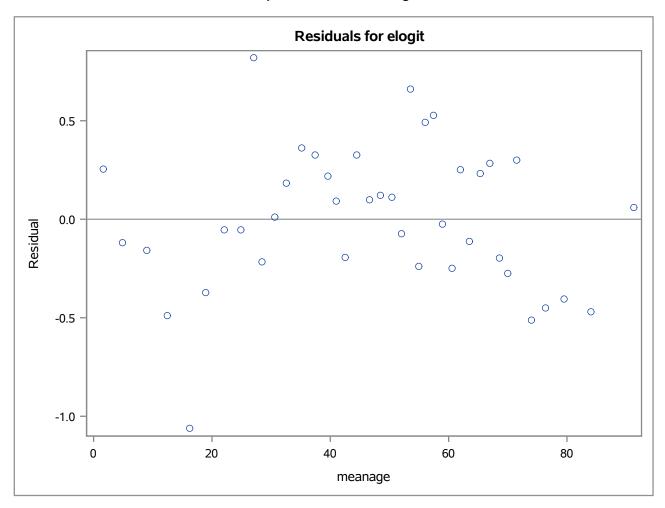
Number of Observations Read	40
Number of Observations Used	40

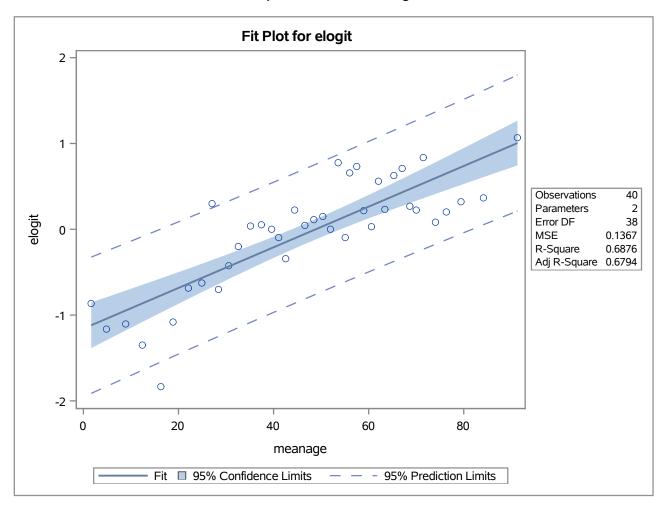
Analysis of Variance						
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F	
Model	1	11.43252	11.43252	83.66	<.0001	
Error	38	5.19311	0.13666			
Corrected Total	39	16.62563				

Root MSE	0.36968	R-Square	0.6876
Dependent Mean	-0.04369	Adj R-Sq	0.6794
Coeff Var	-846.12217		

Parameter Estimates						
Variable DF Parameter Estimate Standard Error t Value Pr > t						
Intercept	1	-1.15766	0.13509	-8.57	<.0001	
meanage	1	0.02369	0.00259	9.15	<.0001	







The CORR Procedure

2 Variables: elogit bin

Simple Statistics							
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum	Label
elogit	40	-0.04369	0.65292	-1.74763	-1.83438	1.06538	
bin	40	19.50000	11.69045	780.00000	0	39.00000	Rank for Variable newage

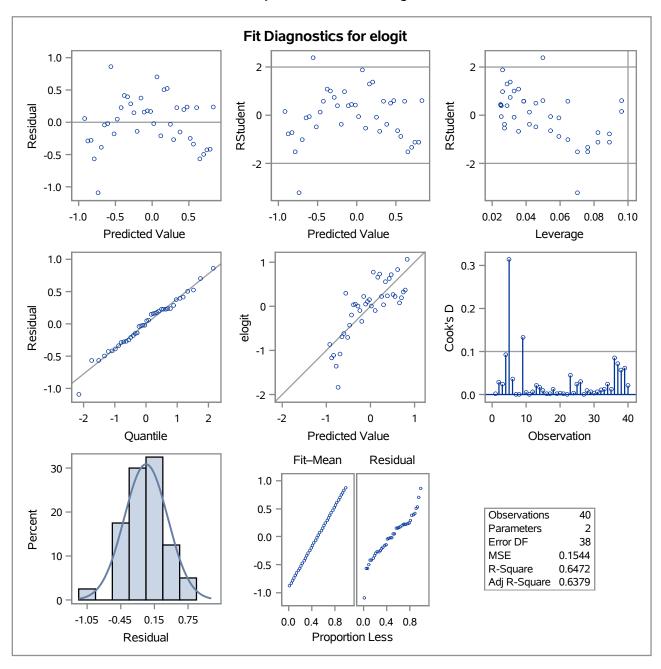
Pearson Correlation Coefficients, N = 40 Prob > r under H0: Rho=0				
	elogit	bin		
elogit	1.00000	0.80448 <.0001		
bin Rank for Variable newage	0.80448 <.0001	1.00000		

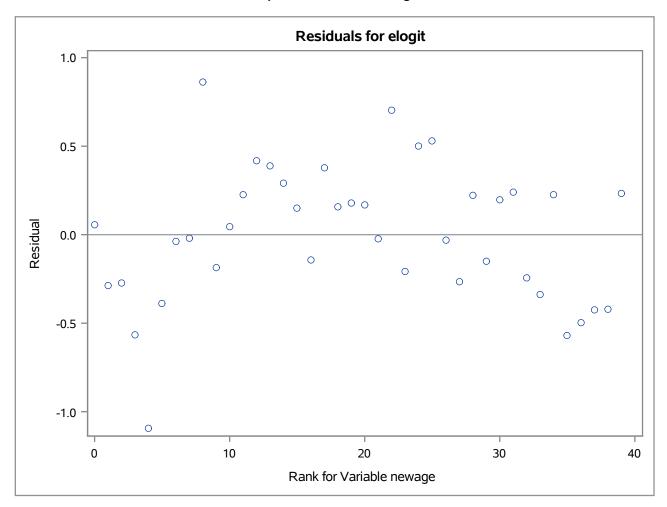
Number of Observations Read	40
Number of Observations Used	40

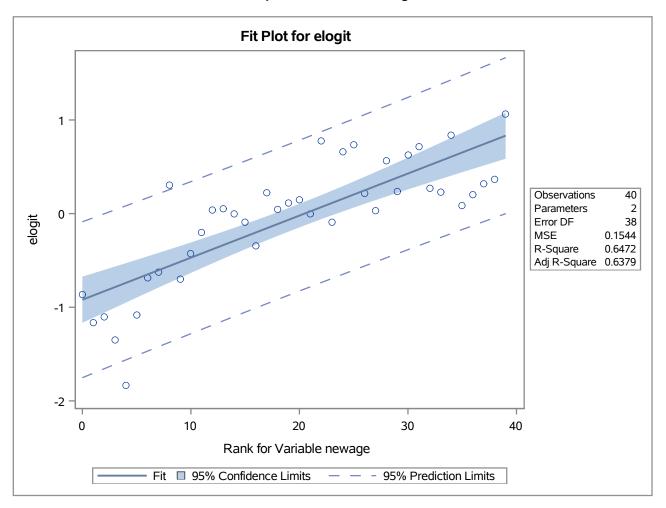
Analysis of Variance						
Source	DF	Sum of Squares	Mean Square	F Value	Pr > F	
Model	1	10.76003	10.76003	69.71	<.0001	
Error	38	5.86560	0.15436			
Corrected Total	39	16.62563				

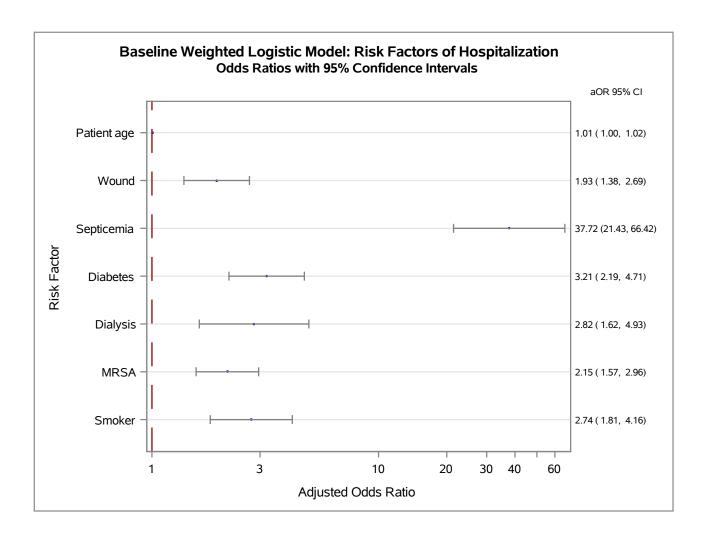
Root MSE	0.39288	R-Square	0.6472
Dependent Mean	-0.04369	Adj R-Sq	0.6379
Coeff Var	-899.23986		

Parameter Estimates						
Variable	ariable Label DF Parameter Standard Error t Value Pr > t					
Intercept	Intercept	1	-0.91984	0.12195	-7.54	<.0001
bin	Rank for Variable newage	1	0.04493	0.00538	8.35	<.0001









Model Information			
Data Set	WORK.COMM		
Response Variable	hosp		
Number of Response Levels	2		
Stratum Variable	INVASIVE		
Number of Strata	2		
Weight Variable	WEIGHT		
Model	Binary Logit		
Optimization Technique	Newton-Raphson		
Variance Adjustment	Degrees of Freedom (DF)		

Variance Estimation				
Method	Taylor Series			
Variance Adjustment	Degrees of Freedom (DF)			

Number of Observations Read	846
Number of Observations Used	846
Sum of Weights Read	2907
Sum of Weights Used	2907

Response Profile					
Ordered Total Total Value hosp Frequency Weigh					
1	No	590	2285.0000		
2	Yes	256	622.0000		

Probability modeled is hosp='Yes'.

Class Level Information				
Value Design Variables				
MRSA	1			
MSSA	0			
No	0			
Yes	1			
No	0			
Yes	1			
No	0			
Yes	1			
	Value MRSA MSSA No Yes No Yes No			

Class Level Information				
Class Value Design Variables				
BSI	No	0		
	Yes	1		
WOUND	No	0		
	Yes	1		

Stratum Information				
Stratum INVASIVE N Obs				
1	No	679		
2	Yes	167		

Model Convergence Status

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

Model Fit Statistics				
Intercept a Criterion Only Covariat				
AIC	3020.422	2305.551		
sc	3026.397	2353.350		
-2 Log L	3018.422	2289.551		

R-Square	0.2218	Max-rescaled R-Square	0.3433	
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Testing Global Null Hypothesis: BETA=0							
Test F Value Num DF Den DF Pr > F							
Likelihood Ratio	30.28	7.0000	5907.98	<.0001			
Score	36.84	7	838	<.0001			
Wald	24.99	7	838	<.0001			

NOTE:

Second-order Rao-Scott design correction 0.0000 applied to the Likelihood Ratio test.

Type 3 Analysis of Effects								
Effect	Effect F Value Num DF Den DF Pr > F							
SMOKER	20.63	1	844	<.0001				
mrsafinal	9.00	1	844	0.0028				
kidney	2.37	1	844	0.1239				
DIABETES	18.60	1	844	<.0001				
BSI	123.15	1	844	<.0001				
WOUND	8.27	1	844	0.0041				
newage	0.65	1	844	0.4201				

Analysis of Maximum Likelihood Estimates					
Parameter	ter Estimate Standard Error t Value Pr > t				
Intercept -2.5522 0.2434 -10.49 <.000				<.0001	
SMOKER	Yes	1.2164	0.2678	4.54	<.0001
mrsafinal	MRSA	0.6836	0.2279	3.00	0.0028
kidney	Yes	0.7472	0.4851	1.54	0.1239
DIABETES Yes 1.219		1.2194	0.2828	4.31	<.0001
BSI Yes 4.5960 0.4142 11.10 <.0001				<.0001	
WOUND Yes 0.6934 0.2410 2.88 0.004				0.0041	
newage 0.00406 0.00504 0.81 0.4201					
NOTE:	The degr	ees of freed	dom for the	t tests is 8	344.

Odds Ratio Estimates				
Effect	Point Estimate	95% Confidence Limits		
SMOKER Yes vs No	3.375	1.995 5.709		
mrsafinal MRSA vs MSSA	1.981	1.266 3.09		
kidney Yes vs No	2.111	0.815 5.471		
DIABETES Yes vs No	3.385	1.943 5.897		
BSI Yes vs No	99.091	43.954	223.391	
WOUND Yes vs No	2.000	1.246	3.211	
newage	1.004	0.994	1.014	

NOTE:

The degrees of freedom in computing the confidence limits is 844.

Association of Predicted Probabilities and Observed Responses						
Percent Concordant 83.3 Somers' D 0.674						
Percent Discordant 15.9 Gamma 0.679						
Percent Tied 0.8 Tau-a 0.285						
Pairs	151040	с	0.837			

Model Information			
Data Set WORK.HOP			
Response Variable	hosp		
Number of Response Levels	2		
Stratum Variable	INVASIVE		
Number of Strata	2		
Weight Variable	WEIGHT		
Model	Binary Logit		
Optimization Technique	Newton-Raphson		
Variance Adjustment	Degrees of Freedom (DF)		

Variance Estimation			
Method Taylor Series			
Variance Adjustment	Degrees of Freedom (DF)		

Number of Observations Read	601
Number of Observations Used	601
Sum of Weights Read	1456
Sum of Weights Used	1456

Response Profile					
Ordered Total Total Value hosp Frequency Weight					
1	No	145	523.00000		
2	Yes	456	933.00000		

Probability modeled is hosp='Yes'.

Class Level Information			
Value	Design Variables		
MRSA	1		
MSSA	0		
No	0		
Yes	1		
No	0		
Yes	1		
No	0		
Yes	1		
	Value MRSA MSSA No Yes No Yes No		

Class Level Information				
Class Value Design Variables				
BSI	No	0		
	Yes	1		
WOUND	No	0		
	Yes	1		

Stratum Information			
Stratum Index	INVASIVE	N Obs	
1	No	274	
2	Yes	327	

Model Convergence Status

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

Model Fit Statistics					
Intercept and Criterion Only Covariates					
AIC	1903.415	1544.926			
sc	1908.698	1587.194			
-2 Log L	1901.415	1528.926			

- Square 0.2257	Max-rescaled R-Square	0.3096	
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Testing Global Null Hypothesis: BETA=0						
Test F Value Num DF Den DF Pr >						
Likelihood Ratio	21.95	6.9999	4192.96	<.0001		
Score	40.62	7	593	<.0001		
Wald	10.58	7	593	<.0001		

NOTE:

Second-order Rao-Scott design correction 0.0000 applied to the Likelihood Ratio test.

Type 3 Analysis of Effects								
Effect	Effect F Value Num DF Den DF Pr > F							
SMOKER	4.50	1	599	0.0342				
mrsafinal	7.76	1	599	0.0055				
kidney	4.68	1	599	0.0309				
DIABETES	12.32	1	599	0.0005				
BSI	36.32	1	599	<.0001				
WOUND	0.17	1	599	0.6819				
newage	1.27	1	599	0.2608				

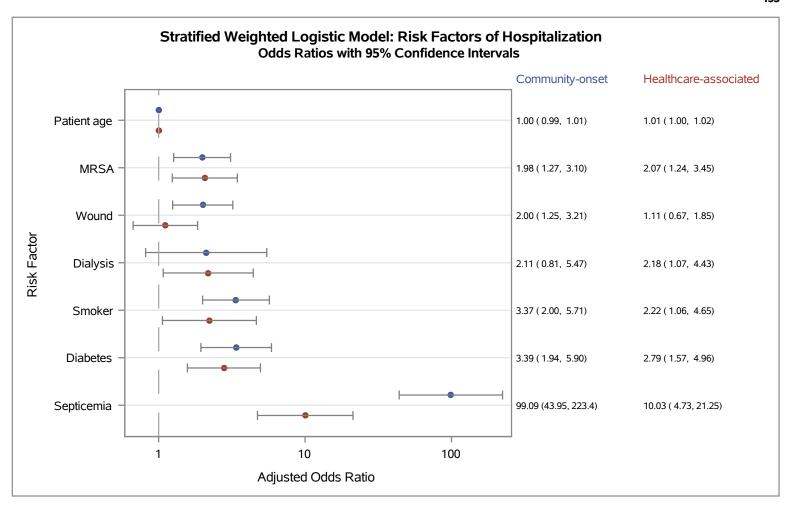
Analysis of Maximum Likelihood Estimates					
Parameter		Estimate	Standard Error	t Value	Pr > t
Intercept		-0.9211	0.3420	-2.69	0.0073
SMOKER	Yes	0.7982	0.3761	2.12	0.0342
mrsafinal	MRSA	0.7257	0.2606	2.79	0.0055
kidney	Yes	0.7799	0.3605	2.16	0.0309
DIABETES	Yes	1.0265	0.2924	3.51	0.0005
BSI	Yes	2.3051	0.3825	6.03	<.0001
WOUND	Yes	0.1059	0.2583	0.41	0.6819
newage		0.00669	0.00594	1.13	0.2608
NOTE: The degrees of freedom for the t tests is 599.					

Odds Ratio Estimates					
Effect	Point Estimate	95% Confidence Limits			
SMOKER Yes vs No	2.222	1.061	4.650		
mrsafinal MRSA vs MSSA	2.066	1.239	3.447		
kidney Yes vs No	2.181	1.075	4.428		
DIABETES Yes vs No	2.791	1.572	4.957		
BSI Yes vs No	10.025	4.730	21.248		
WOUND Yes vs No	1.112	0.669	1.846		
newage	1.007	0.995	1.019		

NOTE:

The degrees of freedom in computing the confidence limits is 599.

Association of Predicted Probabilities and Observed Responses						
Percent Concordant 81.3 Somers' D 0.630						
Percent Discordant	Discordant 18.3 Gamma 0.69					
Percent Tied	0.4 Tau-a 0.2					
Pairs 66120 c 0.815						



The GLIMMIX Procedure

Model Information			
Data Set	S.STAPH		
Response Variable	hosp		
Response Distribution	Binary		
Link Function	Logit		
Variance Function	Default		
Variance Matrix Blocked By	TXHOSP		
Estimation Technique	Maximum Likelihood		
Likelihood Approximation	Gauss-Hermite Quadrature		
Degrees of Freedom Method	Between-Within		
Fixed Effects SE Adjustment	Sandwich - MBN(df,r=1,d=2)		

	Class Level Information				
Class	Levels	Values			
TXHOSP	35	GA002 GA003 GA004 GA006 GA008 GA009 GA010 GA011 GA013 GA015 GA016 GA018 GA020 GA021 GA024 GA026 GA027 GA029 GA030 GA032 GA034 GA040 GA046 GA048 GA050 GA056 GA059 GA065 GA066 GA069 GA070 GA071 GA308 GAMDO OSODC			
mrsafinal	2	MRSA MSSA			
kidney	2	Yes No			
DIABETES	2	Yes No			
SMOKER	2	Yes No			
WOUND	2	Yes No			
BSI	2	Yes No			

Number of Observations Read	1447
Number of Observations Used	1447

Response Profile			
Ordered Value	Total Frequency		
1	No	735	
2	Yes	712	

The GLIMMIX procedure is modeling the probability that hosp='Yes'.

The GLIMMIX Procedure

Dimensions	
G-side Cov. Parameters	1
Columns in X	14
Columns in Z per Subject	1
Subjects (Blocks in V)	35
Max Obs per Subject	447

Optimization Information			
Optimization Technique Dual Quasi-New			
Parameters in Optimization	9		
Lower Boundaries	1		
Upper Boundaries	0		
Fixed Effects	Not Profiled		
Starting From	GLM estimates		
Quadrature Points	1		

	Iteration History					
Iteration	Restarts	Objective Evaluations Function Change		Max Gradient		
0	0	4	2809.7671492		2776.37	
1	0	6	2798.1628166	11.60433259	120.7446	
2	0	4	2738.8833011	59.27951551	40.08431	
3	0	2	2732.1785248	6.70477633	19.49736	
4	0	4	2728.8319698	3.34655496	14.21317	
5	0	2	2728.0560481	0.77592168	4.021616	
6	0	3	2727.5457195	0.51032860	34.32043	
7	0	2	2727.0471694	0.49855013	35.08955	
8	0	2	2726.3982854	0.64888401	8.700023	
9	0	3	2726.1195295	0.27875594	9.737444	
10	0	3	2726.0684443	0.05108520	3.665075	
11	0	3	2726.0633297	0.00511460	1.240708	
12	0	3	2726.0609073	0.00242232	0.580288	
13	0	3	2726.0600159	0.00089147	0.146745	
14	0	3	2726.0600102	0.00000571	0.011663	

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

Fit Statistics		
-2 Log Likelihood	2726.06	
AIC (smaller is better)	2744.06	
AICC (smaller is better)	2744.19	
BIC (smaller is better)	2758.06	
CAIC (smaller is better)	2767.06	
HQIC (smaller is better)	2748.89	

Fit Statistics for Conditional Distribution			
-2 log L(hosp r. effects) 2605.19			
Pearson Chi-Square	1867.72		
Pearson Chi-Square / DF	1.29		

Covariance Parameter Estimates					
Cov Parm Subject Estimate Standard Z Value Pr					Pr > Z
Intercept	TXHOSP	3.3699	2.2234	1.52	0.0648

				Solutions	for Fixed E	Effects					
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t
Intercept							-2.2875	0.5930	34	-3.86	0.0005
SMOKER				Yes			0.5006	0.3209	18	1.56	0.1362
SMOKER				No			0				
mrsafinal	MRSA						0.01493	0.2164	24	0.07	0.9456
mrsafinal	MSSA						0				
kidney		Yes					1.2379	0.3360	22	3.68	0.0013
kidney		No					0				
DIABETES			Yes				0.7665	0.2814	22	2.72	0.0124
DIABETES			No				0				
WOUND					Yes		0.4193	0.2014	21	2.08	0.0498
WOUND					No		0				
BSI						Yes	2.7266	0.3766	24	7.24	<.0001
BSI						No	0				
newage							0.02178	0.005370	4321	4.06	<.0001

The GLIMMIX Procedure

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BS			
			Yes			44.242				No					
MRSA						44.242	MSSA								
	Yes					44.242		No							
		Yes				44.242			No						
				Yes		44.242					No				
					Yes	44.242						No			
						45.242									

Effects of continuous variables are assessed as one unit offsets from the mean.

The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

The GLIMMIX Procedure

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	_newage			
			Yes			44.242				No		44.242			
MRSA						44.242	MSSA					44.242			
	Yes					44.242		No				44.242			
		Yes				44.242			No			44.242			
				Yes		44.242					No	44.242			
					Yes	44.242						44.242			
						45.242						44.242			

The GLIMMIX Procedure

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	Estimate			
			Yes			44.242				No		1.650			
MRSA						44.242	MSSA					1.015			
	Yes					44.242		No				3.448			
		Yes				44.242			No			2.152			
				Yes		44.242					No	1.521			
					Yes	44.242						15.281			
						45.242						1.022			

The GLIMMIX Procedure

						Odds Ra	tio Estimates	i					
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	DF	95% Confi denc e Limit s
			Yes			44.242				No		18	0.841
MRSA						44.242	MSSA					24	0.649
	Yes					44.242		No				22	1.718
		Yes				44.242			No			22	1.201
				Yes		44.242					No	21	1.000
					Yes	44.242						24	7.024
						45.242						4321	1.011

The GLIMMIX Procedure

					C	dds Ratio	Estimates					
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	95% Confi dence Limits
			Yes			44.242				No		3.238
MRSA						44.242	MSSA					1.587
	Yes					44.242		No				6.923
		Yes				44.242			No			3.858
				Yes		44.242					No	2.312
					Yes	44.242						33.242
						45.242						1.033

SMOKER 1 18 2.43 2.43 0.1188 0.1362 mrsafinal 1 24 0.00 0.00 0.9450 0.9456 kidney 1 22 13.57 13.57 0.0002 0.0013 DIABETES 1 22 7.42 7.42 0.0065 0.0124											
Effect			Chi-Square	F Value	Pr > ChiSq	Pr > F					
SMOKER	1	18	2.43	2.43	0.1188	0.1362					
mrsafinal	1	24	0.00	0.00	0.9450	0.9456					
kidney	1	22	13.57	13.57	0.0002	0.0013					
DIABETES	1	22	7.42	7.42	0.0065	0.0124					
WOUND	1	21	4.33	4.33	0.0374	0.0498					
BSI	1	24	52.43	52.43	<.0001	<.0001					
newage	1	4321	16.45	16.45	<.0001	<.0001					

		So	lution for	Randor	n Effects				
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA002	1.0334	0.8120	4355	1.27	0.2032	0.05	-0.5585	2.6253
Intercept	TXHOSP GA003	0.1329	0.6562	4355	0.20	0.8395	0.05	-1.1536	1.4194
Intercept	TXHOSP GA004	1.5172	0.5979	4355	2.54	0.0112	0.05	0.3450	2.6894
Intercept	TXHOSP GA006	0.1466	0.9143	4355	0.16	0.8726	0.05	-1.6460	1.9391
Intercept	TXHOSP GA008	-0.9284	0.6812	4355	-1.36	0.1730	0.05	-2.2638	0.4071
Intercept	TXHOSP GA009	1.6056	0.6720	4355	2.39	0.0169	0.05	0.2882	2.9230
Intercept	TXHOSP GA010	1.1908	0.6076	4355	1.96	0.0501	0.05	-0.00038	2.3820
Intercept	TXHOSP GA011	1.3265	0.5764	4355	2.30	0.0214	0.05	0.1964	2.4566
Intercept	TXHOSP GA013	1.1744	0.7535	4355	1.56	0.1192	0.05	-0.3028	2.6516
Intercept	TXHOSP GA015	1.8955	0.6849	4355	2.77	0.0057	0.05	0.5527	3.2382
Intercept	TXHOSP GA016	0.7896	1.4907	4355	0.53	0.5963	0.05	-2.1328	3.7121

		So	lution for	Randor	n Effects				
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA018	0.5921	0.6871	4355	0.86	0.3889	0.05	-0.7550	1.9392
Intercept	TXHOSP GA020	1.0831	0.5938	4355	1.82	0.0682	0.05	-0.08112	2.2473
Intercept	TXHOSP GA021	1.0662	0.6055	4355	1.76	0.0783	0.05	-0.1208	2.2533
Intercept	TXHOSP GA024	0.7462	0.5985	4355	1.25	0.2126	0.05	-0.4273	1.9196
Intercept	TXHOSP GA026	1.3636	0.6238	4355	2.19	0.0289	0.05	0.1407	2.5865
Intercept	TXHOSP GA027	1.5192	0.6421	4355	2.37	0.0180	0.05	0.2603	2.7781
Intercept	TXHOSP GA029	-0.3851	1.5919	4355	-0.24	0.8089	0.05	-3.5060	2.7358
Intercept	TXHOSP GA030	-1.7577	1.1774	4355	-1.49	0.1356	0.05	-4.0660	0.5506
Intercept	TXHOSP GA032	1.7465	0.6516	4355	2.68	0.0074	0.05	0.4691	3.0239
Intercept	TXHOSP GA034	-0.4074	0.6659	4355	-0.61	0.5407	0.05	-1.7130	0.8981
Intercept	TXHOSP GA040	-4.6288	0.8987	4355	-5.15	<.0001	0.05	-6.3908	-2.8668
Intercept	TXHOSP GA046	0.2668	0.7549	4355	0.35	0.7238	0.05	-1.2133	1.7468
Intercept	TXHOSP GA048	-0.03924	1.0758	4355	-0.04	0.9709	0.05	-2.1483	2.0698
Intercept	TXHOSP GA050	-1.7129	1.4468	4355	-1.18	0.2365	0.05	-4.5494	1.1236
Intercept	TXHOSP GA056	0.3922	1.6040	4355	0.24	0.8068	0.05	-2.7524	3.5368
Intercept	TXHOSP GA059	0.4726	0.6188	4355	0.76	0.4451	0.05	-0.7406	1.6859
Intercept	TXHOSP GA065	-3.4060	1.3883	4355	-2.45	0.0142	0.05	-6.1277	-0.6842
Intercept	TXHOSP GA066	-0.8944	1.3783	4355	-0.65	0.5164	0.05	-3.5965	1.8077
Intercept	TXHOSP GA069	1.4220	1.4749	4355	0.96	0.3350	0.05	-1.4694	4.3135
Intercept	TXHOSP GA070	0.8376	0.6511	4355	1.29	0.1984	0.05	-0.4390	2.1141
Intercept	TXHOSP GA071	-0.1330	0.6855	4355	-0.19	0.8462	0.05	-1.4769	1.2110
Intercept	TXHOSP GA308	-2.3440	1.3530	4355	-1.73	0.0833	0.05	-4.9966	0.3086
Intercept	TXHOSP GAMDO	-3.2892	0.5959	4355	-5.52	<.0001	0.05	-4.4574	-2.1210
Intercept	TXHOSP OSODC	-1.8421	1.4661	4355	-1.26	0.2090	0.05	-4.7164	1.0322

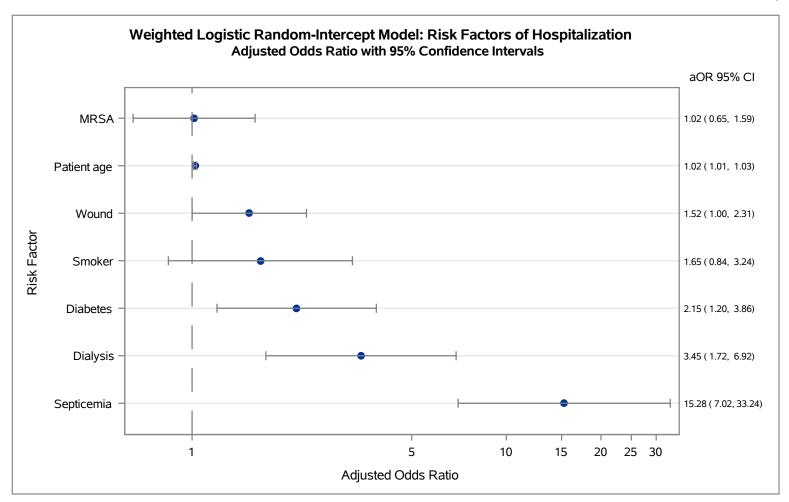
				Empirica	l Correlatio	n Matr	ix for F	ixed Effects	5				
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col1	Col2	Col3	Col4	Col5	Col6
Intercept							1	1.0000	-0.08803		-0.2056		-0.06619
SMOKER				Yes			2	-0.08803	1.0000		0.08159		0.03500
SMOKER				No			3			1.0000			
mrsafinal	MRSA						4	-0.2056	0.08159		1.0000		0.1599
mrsafinal	MSSA						5					1.0000	
kidney		Yes					6	-0.06619	0.03500		0.1599		1.0000
kidney		No					7						
DIABETES			Yes				8	-0.04415	0.2710		0.1505		-0.2076
DIABETES			No				9						
WOUND					Yes		10	-0.1236	0.3643		0.2547		-0.2413
WOUND					No		11						
BSI						Yes	12	-0.07947	0.1116		0.1467		0.02663
BSI						No	13						
newage							14	-0.1829	-0.07769		-0.3066		-0.3054

				Empi	rical Correla	ation M	Matrix fo	r Fixed E	Effects					
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col7	Col8	Col9	Col10	Col11	Col12	Col13
Intercept							1		-0.04415		-0.1236		-0.07947	
SMOKER				Yes			2		0.2710		0.3643		0.1116	
SMOKER				No			3							
mrsafinal	MRSA						4		0.1505		0.2547		0.1467	
mrsafinal	MSSA						5							
kidney		Yes					6		-0.2076		-0.2413		0.02663	
kidney		No					7	1.0000						
DIABETES			Yes				8		1.0000		0.07682		-0.05413	
DIABETES			No				9			1.0000				
WOUND					Yes		10		0.07682		1.0000		0.2771	
WOUND					No		11					1.0000		
BSI						Yes	12		-0.05413		0.2771		1.0000	
BSI						No	13							1.0000
newage							14		-0.1068		0.03849		-0.04147	

Weighted Logistic Random-Intercept Model Conditional on Hospital Cluster

The GLIMMIX Procedure

	Empirical Correlation Matrix for Fixed Effects							
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col14
Intercept							1	-0.1829
SMOKER				Yes			2	-0.07769
SMOKER				No			3	
mrsafinal	MRSA						4	-0.3066
mrsafinal	MSSA						5	
kidney		Yes					6	-0.3054
kidney		No					7	
DIABETES			Yes				8	-0.1068
DIABETES			No				9	
WOUND					Yes		10	0.03849
WOUND					No		11	
BSI						Yes	12	-0.04147
BSI						No	13	
newage							14	1.0000



Model Information						
Data Set	WORK.COMM					
Response Variable	hosp					
Response Distribution	Binary					
Link Function	Logit					
Variance Function	Default					
Variance Matrix Blocked By	TXHOSP					
Estimation Technique	Maximum Likelihood					
Likelihood Approximation	Gauss-Hermite Quadrature					
Degrees of Freedom Method	Between-Within					
Fixed Effects SE Adjustment	Sandwich - MBN(df,r=1,d=2)					

	Class Level Information							
Class	Levels	Values						
TXHOSP	30	GA002 GA003 GA004 GA006 GA008 GA009 GA010 GA011 GA013 GA015 GA016 GA018 GA020 GA021 GA024 GA026 GA027 GA030 GA032 GA034 GA040 GA046 GA048 GA050 GA059 GA066 GA069 GA070 GA071 GAMDO						
mrsafinal	2	MRSA MSSA						
kidney	2	Yes No						
DIABETES	2	Yes No						
SMOKER	2	Yes No						
WOUND	2	Yes No						
BSI	2	Yes No						

Number of Observations Read	846	
Number of Observations Used	846	

Response Profile								
Ordered Value	hosp	Total Frequency						
1	No	590						
2	Yes	256						

The GLIMMIX procedure is modeling the probability that hosp='Yes'.

Dimensions					
G-side Cov. Parameters	1				
Columns in X	14				
Columns in Z per Subject	1				
Subjects (Blocks in V)	30				
Max Obs per Subject	380				

Optimization Information						
Optimization Technique	Dual Quasi-Newton					
Parameters in Optimization	9					
Lower Boundaries	1					
Upper Boundaries	0					
Fixed Effects	Not Profiled					
Starting From	GLM estimates					
Quadrature Points	1					

	Iteration History								
Iteration	Restarts	Evaluations	Objective Function Change		Max Gradient				
0	0	4	1501.0917107		2753.628				
1	0	6	1480.828851	20.26285972	79.96838				
2	0	2	1446.3052458	34.52360519	25.9747				
3	0	2	1438.6102748	7.69497099	25.24847				
4	0	4	1434.7206514	3.88962333	47.57425				
5	0	4	1431.4966568	3.22399465	12.60808				
6	0	3	1431.0656799	0.43097691	13.71226				
7	0	4	1430.0803129	0.98536704	5.962923				
8	0	3	1429.733555	0.34675782	14.80668				
9	0	3	1429.6672263	0.06632873	2.606062				
10	0	3	1429.6524119	0.01481443	1.815746				
11	0	2	1429.6338936	0.01851827	2.527292				
12	0	3	1429.6301493	0.00374429	0.276731				
13	0	3	1429.6300976	0.00005169	0.184168				
14	0	3	1429.6300848	0.00001285	0.016909				

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

Fit Statistics						
-2 Log Likelihood	1429.63					
AIC (smaller is better)	1447.63					
AICC (smaller is better)	1447.85					
BIC (smaller is better)	1460.24					
CAIC (smaller is better)	1469.24					
HQIC (smaller is better)	1451.66					

Fit Statistics for Conditional Distribution						
-2 log L(hosp r. effects) 1329.79						
Pearson Chi-Square	1214.52					
Pearson Chi-Square / DF	1.44					

Covariance Parameter Estimates									
Cov Parm	Subject	Estimate	Standard Error	Z Value	Pr > Z				
Intercept	TXHOSP	4.1805	2.3097	1.81	0.0351				

	Solutions for Fixed Effects										
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t
Intercept							-2.5290	0.6961	29	-3.63	0.0011
SMOKER				Yes			0.6718	0.4220	17	1.59	0.1298
SMOKER				No			0				
mrsafinal	MRSA						-0.1810	0.2851	22	-0.64	0.5320
mrsafinal	MSSA						0				
kidney		Yes					1.6130	0.6778	10	2.38	0.0386
kidney		No					0				
DIABETES			Yes				1.1581	0.4643	17	2.49	0.0232
DIABETES			No				0				
WOUND					Yes		0.5562	0.3572	16	1.56	0.1391
WOUND					No		0				
BSI						Yes	4.0945	0.7998	19	5.12	<.0001
BSI						No	0				
newage							0.02049	0.007379	2870	2.78	0.0055

	Odds Ratio Estimates											
mrsafinal kidney DIABETES SMOKER WOUND BSI newage _mrsafinal _kidney _DIABETES _								_SMOKER	_WOUND	_BSI		
			Yes			39.545				No		
MRSA						39.545	MSSA					
	Yes					39.545		No				
		Yes				39.545			No			
				Yes		39.545					No	
					Yes	39.545						No
						40.545						

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates											
mrsafinal kidney DIABETES SMOKER WOUND BSI newage _mrsafinal _kidney _DIABETES									_SMOKER	_wound	_newage	
			Yes			39.545				No		39.545
MRSA						39.545	MSSA					39.545
	Yes					39.545		No				39.545
		Yes				39.545			No			39.545
				Yes		39.545					No	39.545
					Yes	39.545						39.545
						40.545						39.545

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates											
mrsafinal kidney DIABETES SMOKER WOUND BSI newage _mrsafinal _kidney _DIABETES								_DIABETES	_SMOKER	_WOUND	Estimate	
			Yes			39.545				No		1.958
MRSA						39.545	MSSA					0.834
	Yes					39.545		No				5.018
		Yes				39.545			No			3.184
				Yes		39.545					No	1.744
					Yes	39.545						60.007
						40.545						1.021

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates												
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	DF	95% Confi dence Limits
			Yes			39.545				No		17	0.804
MRSA						39.545	MSSA					22	0.462
	Yes					39.545		No				10	1.108
		Yes				39.545			No			17	1.196
				Yes		39.545					No	16	0.818
					Yes	39.545						19	11.252
						40.545						2870	1.006

Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

	Odds Ratio Estimates											
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	95% Confid ence Limits
			Yes			39.545				No		4.768
MRSA						39.545	MSSA					1.507
	Yes					39.545		No				22.721
		Yes				39.545			No			8.480
				Yes		39.545					No	3.719
					Yes	39.545						320.033
						40.545						1.036

Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

Type III Tests of Fixed Effects							
Effect	Num DF	Den DF	F Value	Pr > F			
SMOKER	1	17	2.53	0.1298			
mrsafinal	1	22	0.40	0.5320			
kidney	1	10	5.66	0.0386			
DIABETES	1	17	6.22	0.0232			
WOUND	1	16	2.42	0.1391			
BSI	1	19	26.21	<.0001			
newage	1	2870	7.71	0.0055			

	Solution for Random Effects								
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA002	-1.3852	1.0465	2899	-1.32	0.1857	0.05	-3.4372	0.6668
Intercept	TXHOSP GA003	-0.6542	0.8484	2899	-0.77	0.4407	0.05	-2.3177	1.0093
Intercept	TXHOSP GA004	0.7617	0.6833	2899	1.11	0.2651	0.05	-0.5781	2.1015
Intercept	TXHOSP GA006	-1.3408	1.4341	2899	-0.93	0.3499	0.05	-4.1527	1.4712
Intercept	TXHOSP GA008	-1.6991	0.9333	2899	-1.82	0.0688	0.05	-3.5291	0.1309
Intercept	TXHOSP GA009	1.3989	0.8269	2899	1.69	0.0908	0.05	-0.2224	3.0202
Intercept	TXHOSP GA010	1.2832	0.6953	2899	1.85	0.0651	0.05	-0.08013	2.6465
Intercept	TXHOSP GA011	1.2810	0.6697	2899	1.91	0.0559	0.05	-0.03207	2.5941
Intercept	TXHOSP GA013	2.4816	0.8876	2899	2.80	0.0052	0.05	0.7413	4.2219
Intercept	TXHOSP GA015	2.3732	0.8225	2899	2.89	0.0039	0.05	0.7605	3.9860
Intercept	TXHOSP GA016	0.5821	1.6877	2899	0.34	0.7302	0.05	-2.7270	3.8912
Intercept	TXHOSP GA018	1.0116	0.8259	2899	1.22	0.2207	0.05	-0.6077	2.6310

	Solution for Random Effects									
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper	
Intercept	TXHOSP GA020	0.9734	0.6947	2899	1.40	0.1613	0.05	-0.3887	2.3355	
Intercept	TXHOSP GA021	0.6896	0.7048	2899	0.98	0.3279	0.05	-0.6923	2.0715	
Intercept	TXHOSP GA024	-0.09621	0.7157	2899	-0.13	0.8931	0.05	-1.4995	1.3071	
Intercept	TXHOSP GA026	1.4177	0.7398	2899	1.92	0.0554	0.05	-0.03294	2.8683	
Intercept	TXHOSP GA027	1.7194	0.7319	2899	2.35	0.0189	0.05	0.2842	3.1545	
Intercept	TXHOSP GA030	-1.7086	1.3302	2899	-1.28	0.1991	0.05	-4.3168	0.8996	
Intercept	TXHOSP GA032	1.5642	0.7456	2899	2.10	0.0360	0.05	0.1022	3.0262	
Intercept	TXHOSP GA034	-1.4768	0.8780	2899	-1.68	0.0927	0.05	-3.1983	0.2447	
Intercept	TXHOSP GA040	-4.0165	1.0574	2899	-3.80	0.0001	0.05	-6.0899	-1.9432	
Intercept	TXHOSP GA046	2.8069	1.3843	2899	2.03	0.0427	0.05	0.09259	5.5212	
Intercept	TXHOSP GA048	-0.7364	1.5905	2899	-0.46	0.6434	0.05	-3.8551	2.3823	
Intercept	TXHOSP GA050	-2.4485	1.6212	2899	-1.51	0.1311	0.05	-5.6272	0.7302	
Intercept	TXHOSP GA059	0.8361	0.7259	2899	1.15	0.2495	0.05	-0.5872	2.2595	
Intercept	TXHOSP GA066	-0.7882	1.5712	2899	-0.50	0.6160	0.05	-3.8690	2.2927	
Intercept	TXHOSP GA069	0.3367	1.8053	2899	0.19	0.8520	0.05	-3.2030	3.8764	
Intercept	TXHOSP GA070	1.3729	0.8034	2899	1.71	0.0876	0.05	-0.2024	2.9483	
Intercept	TXHOSP GA071	-1.0502	0.8703	2899	-1.21	0.2276	0.05	-2.7566	0.6561	
Intercept	TXHOSP GAMDO	-4.2320	0.7083	2899	-5.97	<.0001	0.05	-5.6209	-2.8431	

Model Info	ormation
Data Set	WORK.HOP
Response Variable	hosp
Response Distribution	Binary
Link Function	Logit
Variance Function	Default
Variance Matrix Blocked By	TXHOSP
Estimation Technique	Maximum Likelihood
Likelihood Approximation	Gauss-Hermite Quadrature
Degrees of Freedom Method	Between-Within
Fixed Effects SE Adjustment	Sandwich - MBN(df,r=1,d=2)

	Class Level Information							
Class	Levels	Values						
TXHOSP	31	GA002 GA003 GA004 GA006 GA008 GA009 GA010 GA011 GA013 GA015 GA018 GA020 GA021 GA024 GA026 GA027 GA029 GA032 GA034 GA040 GA046 GA048 GA056 GA059 GA065 GA069 GA070 GA071 GA308 GAMDO OSODC						
mrsafinal	2	MRSA MSSA						
kidney	2	Yes No						
DIABETES	2	Yes No						
SMOKER	2	Yes No						
WOUND	2	Yes No						
BSI	2	Yes No						

Number of Observations Read	601
Number of Observations Used	601

Response Profile					
Ordered Value	hosp	Total Frequency			
1	No	145			
2	Yes	456			

The GLIMMIX procedure is modeling the probability that hosp='Yes'.

Dimensions	
G-side Cov. Parameters	1
Columns in X	14
Columns in Z per Subject	1
Subjects (Blocks in V)	31
Max Obs per Subject	120

Optimization Information										
Optimization Technique	Dual Quasi-Newton									
Parameters in Optimization	9									
Lower Boundaries	1									
Upper Boundaries	0									
Fixed Effects	Not Profiled									
Starting From	GLM estimates									
Quadrature Points	5									

	Iteration History												
Iteration	Restarts	Evaluations	Objective Function	Change	Max Gradient								
0	0	4	1146.8534811		326.408								
1	0	5	1146.5142917	0.33918935	22.92349								
2	0	4	1135.1806641	11.33362762	8.293453								
3	0	2	1132.1465315	3.03413262	9.172207								
4	0	4	1130.8451183	1.30141320	15.36075								
5	0	4	1129.7493425	1.09577577	2.771987								
6	0	2	1129.3311559	0.41818661	59.56893								
7	0	2	1128.638522	0.69263391	3.230332								
8	0	3	1128.3265204	0.31200159	26.83417								
9	0	2	1128.1767489	0.14977150	41.32122								
10	0	2	1128.127342	0.04940697	49.8322								
11	0	2	1128.0514506	0.07589132	19.01861								
12	0	3	1128.0280862	0.02336444	5.386648								
13	0	3	1128.0140478	0.01403844	0.498978								
14	0	3	1128.0139151	0.00013265	0.16206								
15	0	3	1128.0139063	0.00000880	0.149955								

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

Fit Statistics	
-2 Log Likelihood	1128.01
AIC (smaller is better)	1146.01
AICC (smaller is better)	1146.32
BIC (smaller is better)	1158.92
CAIC (smaller is better)	1167.92
HQIC (smaller is better)	1150.22

Fit Statistics for Condi Distribution	tional
-2 log L(hosp r. effects)	1034.77
Pearson Chi-Square	596.07
Pearson Chi-Square / DF	0.99

	Covaria	nce Parame	eter Estimate	es	
Cov Parm	Subject	Estimate	Standard Error	Z Value	Pr > Z
Intercept	TXHOSP	4.0459	3.0019	1.35	0.0889

	Solutions for Fixed Effects													
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t			
Intercept							-0.4119	0.7588	30	-0.54	0.5912			
SMOKER				Yes			0.4916	0.7261	16	0.68	0.5081			
SMOKER				No			0							
mrsafinal	MRSA						0.2375	0.4195	22	0.57	0.5770			
mrsafinal	MSSA						0							
kidney		Yes					0.8999	0.4452	19	2.02	0.0576			
kidney		No					0							
DIABETES			Yes				0.6596	0.3741	19	1.76	0.0940			
DIABETES			No				0							
WOUND					Yes		-0.1607	0.3984	19	-0.40	0.6912			
WOUND					No		0							
BSI						Yes	1.6658	0.5673	21	2.94	0.0079			
BSI						No	0							
newage							0.005249	0.007870	1418	0.67	0.5049			

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BSI			
			Yes			53.62				No					
MRSA						53.62	MSSA								
	Yes					53.62		No							
		Yes				53.62			No						
				Yes		53.62					No				
					Yes	53.62						No			
						54.62									

Effects of continuous variables are assessed as one unit offsets from the mean.

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_newage			
			Yes			53.62				No		53.62			
MRSA						53.62	MSSA					53.62			
	Yes					53.62		No				53.62			
		Yes				53.62			No			53.62			
				Yes		53.62					No	53.62			
					Yes	53.62						53.62			
						54.62						53.62			

Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

	Odds Ratio Estimates														
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_wound	Estimate			
			Yes			53.62				No		1.635			
MRSA						53.62	MSSA					1.268			
	Yes					53.62		No				2.459			
		Yes				53.62			No			1.934			
				Yes		53.62					No	0.852			
					Yes	53.62						5.290			
						54.62						1.005			

Effects of continuous variables are assessed as one unit offsets from the mean.

						Odds Ra	tio Estimates	;					
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	DF	95% Confi denc e Limit s
			Yes			53.62				No		16	0.351
MRSA						53.62	MSSA					22	0.531
	Yes					53.62		No				19	0.969
		Yes				53.62			No			19	0.884
				Yes		53.62					No	19	0.370
					Yes	53.62						21	1.626
						54.62						1418	0.990

Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

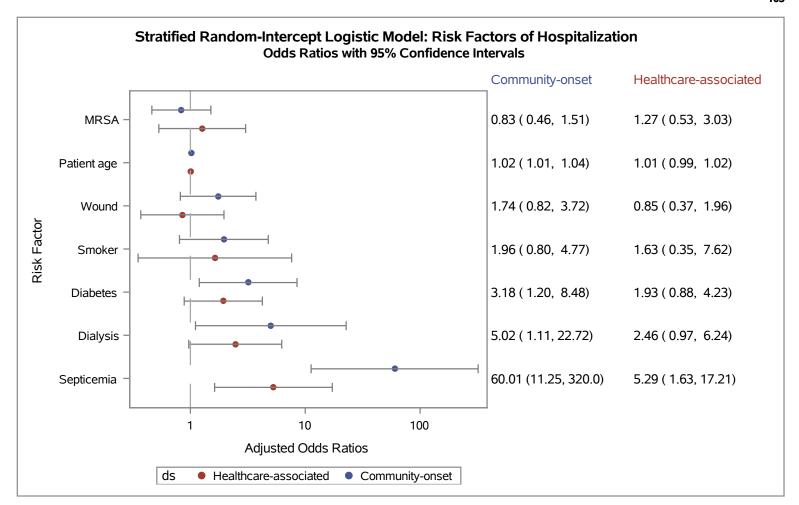
					С	dds Ratio	Estimates					
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	95% Confi dence Limits
			Yes			53.62				No		7.620
MRSA						53.62	MSSA					3.027
	Yes					53.62		No				6.245
		Yes				53.62			No			4.232
				Yes		53.62					No	1.960
					Yes	53.62						17.212
						54.62						1.021

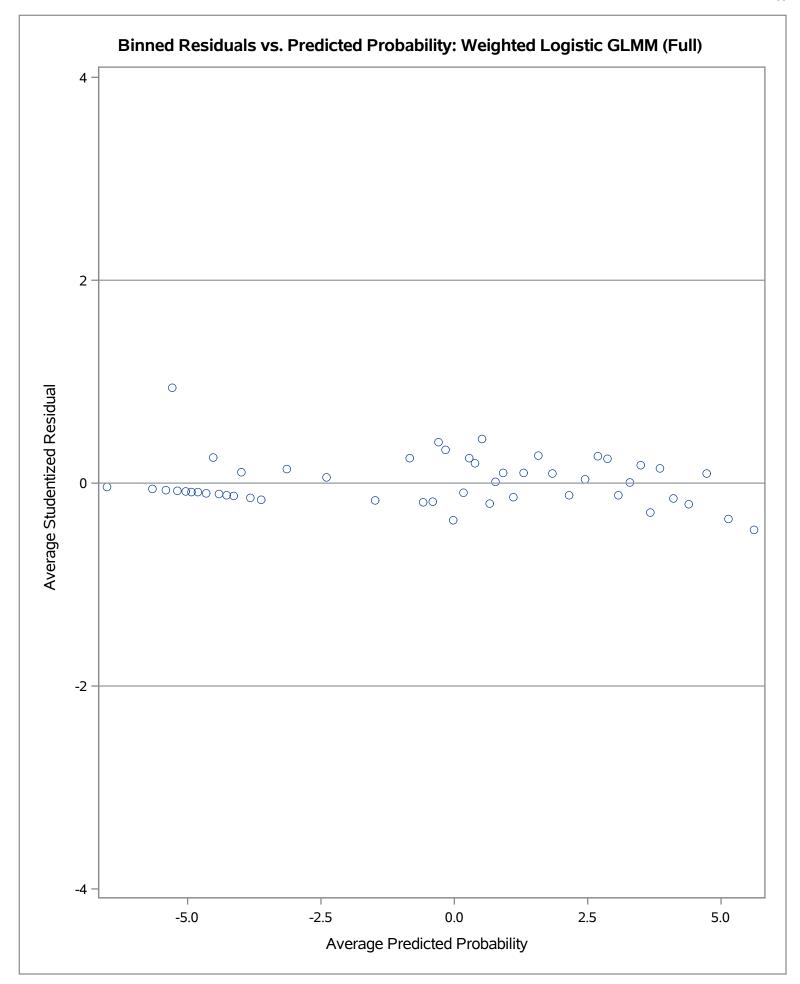
Effects of continuous variables are assessed as one unit offsets from the mean. The AT suboption modifies the reference value and the UNIT suboption modifies the offsets.

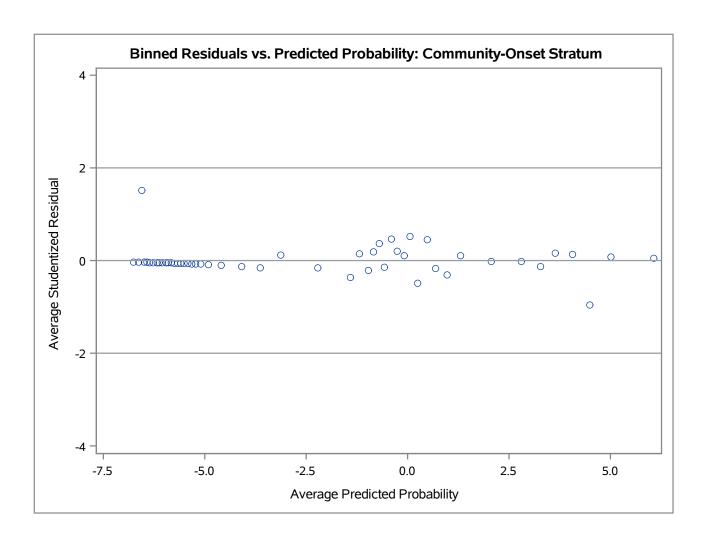
Type III Tests of Fixed Effects							
Effect	Num DF	Den DF	F Value	Pr > F			
SMOKER	1	16	0.46	0.5081			
mrsafinal	1	22	0.32	0.5770			
kidney	1	19	4.09	0.0576			
DIABETES	1	19	3.11	0.0940			
WOUND	1	19	0.16	0.6912			
BSI	1	21	8.62	0.0079			
newage	1	1418	0.44	0.5049			

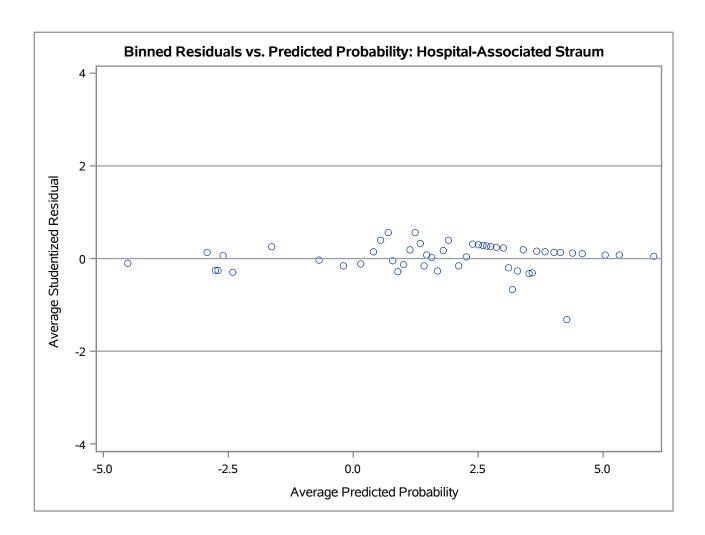
	Solution for Random Effects								
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA002	2.6027	1.3808	1448	1.88	0.0596	0.05	-0.1059	5.3113
Intercept	TXHOSP GA003	0.5977	0.8245	1448	0.72	0.4686	0.05	-1.0196	2.2150
Intercept	TXHOSP GA004	1.8489	0.7062	1448	2.62	0.0089	0.05	0.4637	3.2342
Intercept	TXHOSP GA006	1.3323	1.5733	1448	0.85	0.3972	0.05	-1.7539	4.4186
Intercept	TXHOSP GA008	-0.9807	0.8354	1448	-1.17	0.2406	0.05	-2.6195	0.6580
Intercept	TXHOSP GA009	1.0063	0.7729	1448	1.30	0.1932	0.05	-0.5099	2.5224
Intercept	TXHOSP GA010	0.8390	0.7660	1448	1.10	0.2736	0.05	-0.6636	2.3417
Intercept	TXHOSP GA011	0.8599	0.6431	1448	1.34	0.1814	0.05	-0.4016	2.1213
Intercept	TXHOSP GA013	-2.0640	1.1511	1448	-1.79	0.0732	0.05	-4.3219	0.1940
Intercept	TXHOSP GA015	-0.1339	0.8388	1448	-0.16	0.8732	0.05	-1.7793	1.5115
Intercept	TXHOSP GA018	0.1910	0.8579	1448	0.22	0.8238	0.05	-1.4918	1.8738
Intercept	TXHOSP GA020	1.6896	0.7831	1448	2.16	0.0311	0.05	0.1534	3.2258

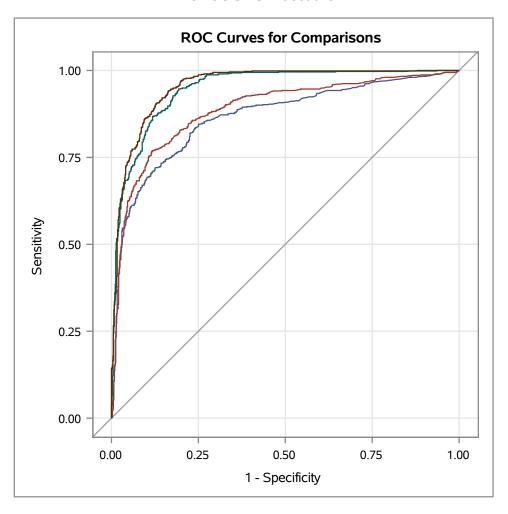
	Solution for Random Effects								
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA021	1.0536	0.6838	1448	1.54	0.1236	0.05	-0.2877	2.3949
Intercept	TXHOSP GA024	0.5776	0.6655	1448	0.87	0.3856	0.05	-0.7279	1.8831
Intercept	TXHOSP GA026	0.9917	0.7209	1448	1.38	0.1691	0.05	-0.4224	2.4058
Intercept	TXHOSP GA027	0.02234	0.8926	1448	0.03	0.9800	0.05	-1.7286	1.7733
Intercept	TXHOSP GA029	-1.0374	1.5837	1448	-0.66	0.5126	0.05	-4.1440	2.0692
Intercept	TXHOSP GA032	2.9202	1.3270	1448	2.20	0.0279	0.05	0.3170	5.5233
Intercept	TXHOSP GA034	-0.3605	0.7015	1448	-0.51	0.6074	0.05	-1.7366	1.0156
Intercept	TXHOSP GA040	-4.5150	1.1279	1448	-4.00	<.0001	0.05	-6.7275	-2.3025
Intercept	TXHOSP GA046	-1.2222	0.8766	1448	-1.39	0.1634	0.05	-2.9417	0.4972
Intercept	TXHOSP GA048	0.7424	1.6589	1448	0.45	0.6546	0.05	-2.5117	3.9965
Intercept	TXHOSP GA056	0.5880	1.6793	1448	0.35	0.7263	0.05	-2.7061	3.8821
Intercept	TXHOSP GA059	-0.07575	0.7352	1448	-0.10	0.9180	0.05	-1.5179	1.3664
Intercept	TXHOSP GA065	-3.3498	1.5351	1448	-2.18	0.0293	0.05	-6.3611	-0.3385
Intercept	TXHOSP GA069	1.0688	1.6253	1448	0.66	0.5109	0.05	-2.1194	4.2571
Intercept	TXHOSP GA070	0.1242	0.7144	1448	0.17	0.8620	0.05	-1.2772	1.5255
Intercept	TXHOSP GA071	0.3750	0.9252	1448	0.41	0.6853	0.05	-1.4400	2.1900
Intercept	TXHOSP GA308	-2.6334	1.4665	1448	-1.80	0.0727	0.05	-5.5100	0.2432
Intercept	TXHOSP GAMDO	-2.4710	0.6609	1448	-3.74	0.0002	0.05	-3.7674	-1.1746
Intercept	TXHOSP OSODC	-1.8292	1.5960	1448	-1.15	0.2519	0.05	-4.9600	1.3016











ROC Association Statistics								
		Mann-Whitney						
ROC Model	Area	Standard Error	95% Wald Confidence Limits		Somers' D Gamma	Gamma	Tau-a	
Baseline_Weighted_Logistic_Model	0.8654	0.00980	0.8462	0.8846	0.7309	0.7314	0.3656	
Stratified Weighted Logistic Model	0.8869	0.00900	0.8693	0.9046	0.7739	0.7742	0.3871	
Random-effects Weighted Logistic Model	0.9471	0.00553	0.9362	0.9579	0.8942	0.8942	0.4473	
Stratified, Random-effects Weighted Logistic Model	0.9557	0.00502	0.9459	0.9656	0.9115	0.9115	0.4559	

Model Information				
Data Set	WORK.TRAINING			
Response Variable	HOSPITAL			
Number of Response Levels	2			
Weight Variable	WEIGHT			
Model	binary logit			
Optimization Technique	Fisher's scoring			

Number of Observations Read	1155
Number of Observations Used	1155
Sum of Weights Read	3510
Sum of Weights Used	3510

Response Profile						
Ordered Value	HOSPITAL	Total Frequency	Total Weight			
1	0	567	2169.0000			
2	1	588	1341.0000			

Probability modeled is HOSPITAL='1'.

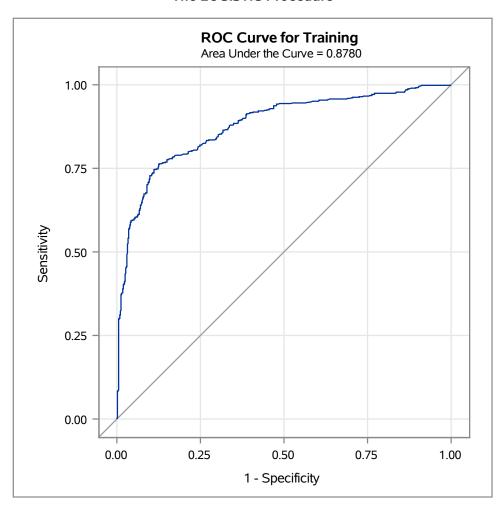
Model Convergence Status
Convergence criterion (GCONV=1E-8) satisfied.

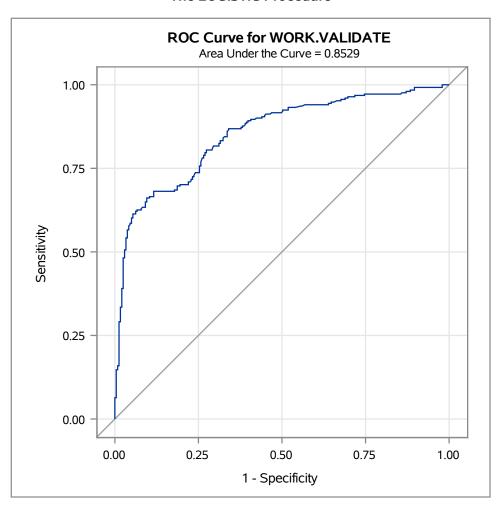
Model Fit Statistics						
Criterion	Intercept Only	Intercept and Covariates				
AIC	4670.717	3330.289				
sc	4675.769	3375.756				
-2 Log L	4668.717	3312.289				

Testing Global Null Hypothesis: BETA=0						
Test Chi-Square DF Pr > ChiS						
Likelihood Ratio	1356.4278	8	<.0001			
Score	1147.9990	8	<.0001			
Wald	685.2433	8	<.0001			

	Analysis of Maximum Likelihood Estimates								
Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq				
Intercept	1	-1.1101	0.1331	69.5132	<.0001				
UND	1	-1.0933	0.1128	93.9653	<.0001				
DISLTC8	1	2.0516	0.2517	66.4630	<.0001				
DIABETES	1	0.8131	0.1074	57.3384	<.0001				
RENAL	1	1.2007	0.1559	59.3029	<.0001				
SST	1	-0.7378	0.0919	64.4613	<.0001				
mrsafinal	1	0.8302	0.0919	81.6576	<.0001				
newage	1	0.00818	0.00212	14.8212	0.0001				
BSI	1	2.7088	0.2573	110.8282	<.0001				

Odds Ratio Estimates						
Effect	Point Estimate	95% Wald Confidence Limits				
UND	0.335	0.269	0.418			
DISLTC8	7.780	4.751	12.741			
DIABETES	2.255	1.827	2.783			
RENAL	3.322	2.448	4.510			
SST	0.478	0.399	0.573			
mrsafinal	2.294	1.916	2.746			
newage	1.008	1.004	1.012			
BSI	15.012	9.066	24.857			





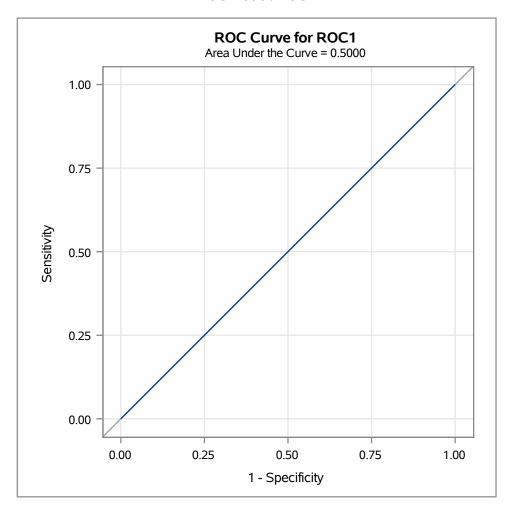
ROC Model: ROC1

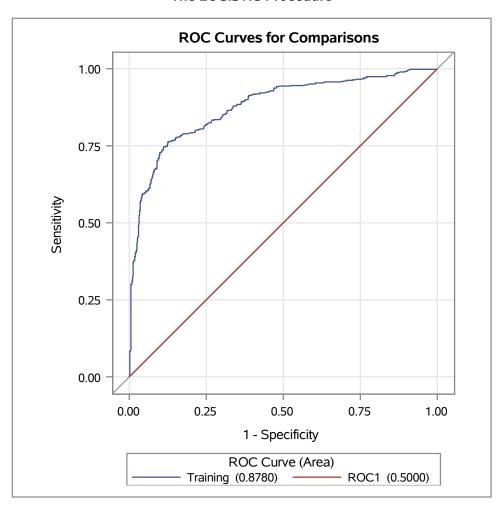
Model Convergence Status Convergence criterion (GCONV=1E-8) satisfied.

> -2 Log L 4668.717

Analysis of Maximum Likelihood Estimates						
Parameter DF Estimate Standard Wald Chi-Square Pr > ChiSq						
Intercept	1	-0.4809	0.0347	191.6027	<.0001	

ROC Model: ROC1





ROC Association Statistics								
	Mann-Whitney							
ROC Model	Area	Standard Error	95% Wald Confidence Limits		Somers' D	Gamma	Tau-a	
Training	0.8780	0.0102	0.8581	0.8980	0.7561	0.7563	0.3782	
ROC1	0.5000	0	0.5000	0.5000	0		0	

ROC Contrast Test Results						
Contrast DF Chi-Square Pr > ChiSc						
Reference = Training	1	1378.7261	<.0001			

Model Information					
Data Set	WORK.TRAINING				
Response Variable	HOSPITAL				
Number of Response Levels	2				
Weight Variable	WEIGHT				
Model	binary logit				
Optimization Technique	Fisher's scoring				

Number of Observations Read	1155
Number of Observations Used	1155
Sum of Weights Read	3510
Sum of Weights Used	3510

Response Profile						
Ordered Value	HOSPITAL	Total Frequency	Total Weight			
1	0	567	2169.0000			
2	1	588	1341.0000			

Probability modeled is HOSPITAL='1'.

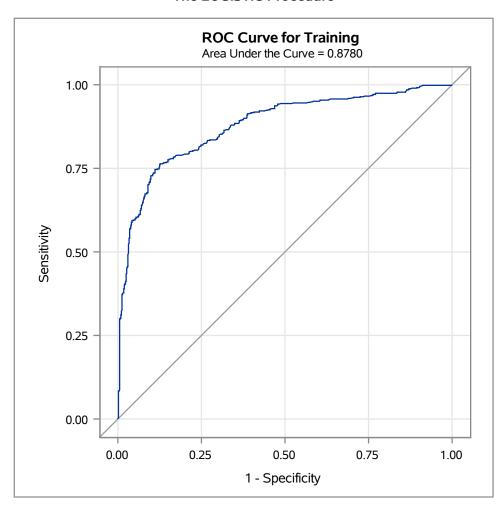
Model Convergence Status
Convergence criterion (GCONV=1E-8) satisfied.

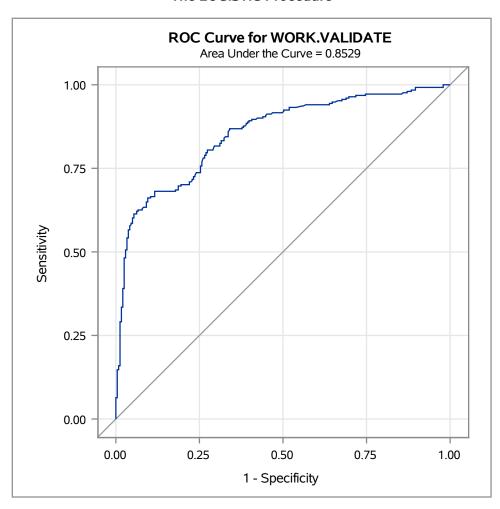
Model Fit Statistics						
Criterion	Intercept Only	Intercept and Covariates				
AIC	4670.717	3330.289				
sc	4675.769	3375.756				
-2 Log L	4668.717	3312.289				

Testing Global Null Hypothesis: BETA=0						
Test Chi-Square DF Pr > Chi						
Likelihood Ratio	1356.4278	8	<.0001			
Score	1147.9990	8	<.0001			
Wald	685.2433	8	<.0001			

Analysis of Maximum Likelihood Estimates							
Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq		
Intercept	1	-1.1101	0.1331	69.5132	<.0001		
UND	1	-1.0933	0.1128	93.9653	<.0001		
DISLTC8	1	2.0516	0.2517	66.4630	<.0001		
DIABETES	1	0.8131	0.1074	57.3384	<.0001		
RENAL	1	1.2007	0.1559	59.3029	<.0001		
SST	1	-0.7378	0.0919	64.4613	<.0001		
mrsafinal	1	0.8302	0.0919	81.6576	<.0001		
newage	1	0.00818	0.00212	14.8212	0.0001		
BSI	1	2.7088	0.2573	110.8282	<.0001		

Odds Ratio Estimates						
Effect	Point Estimate	95% Wald Confidence Limits				
UND	0.335	0.269	0.418			
DISLTC8	7.780	4.751	12.741			
DIABETES	2.255	1.827	2.783			
RENAL	3.322	2.448	4.510			
SST	0.478	0.399 0.5				
mrsafinal	2.294	1.916 2.7				
newage	1.008	1.004	1.012			
BSI	15.012	9.066	24.857			





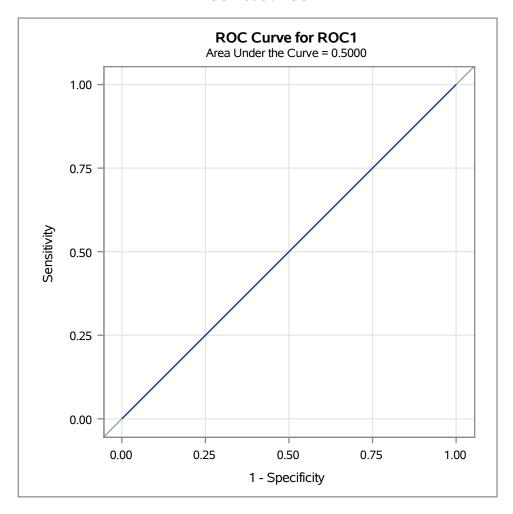
ROC Model: ROC1

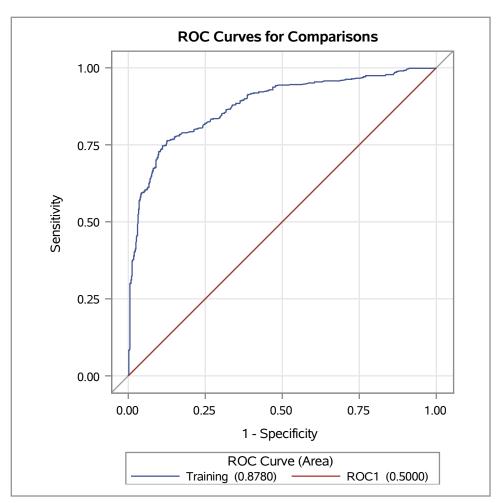
Model Convergence Status Convergence criterion (GCONV=1E-8) satisfied.

> 4668.717 -2 Log L

Analysis of Maximum Likelihood Estimates						
Parameter DF Estimate Standard Wald Chi-Square Pr > ChiSq						
Intercept	1	-0.4809	0.0347	191.6027	<.0001	

ROC Model: ROC1





ROC Association Statistics									
	Mann-Whitney								
ROC Model	Area	Standard Error	95% Wald Confidence Limits		Somers' D	Gamma	Tau-a		
Training	0.8780	0.0102	0.8581	0.8980	0.7561	0.7563	0.3782		
ROC1	0.5000	0	0.5000	0.5000	0		0		

ROC Contrast Test Results							
Contrast	DF	Chi-Square	Pr > ChiSq				
Reference = Training	1	1378.7261	<.0001				