SAS Modeling Output Baseline Weighted Logistic Regression Model

The SURVEYLOGISTIC Procedure

Model Information			
Data Set	S.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'.

Clas	s Level I	nformation
Class	Value	Design Variables
mrsafinal	MRSA	1
	MSSA	(
kidney	No	(
	Yes	1
DIABETES	No	
	Yes	
SMOKER	No	
	Yes	
BSI	No	
	Yes	1
WOUND	No	
	Yes	

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

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

Model Fit Statistics			
Criterion	Intercept Only	Intercept and 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

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	

Type 3 Analysis of Effects					
Effect	F Value	Num DF	Den DF	Pr>	
SMOKER	22.54	1	1445	<.000	
mrsafinal	22.41	1	1445	<.000	
kidney	13.38	1	1445	0.0003	

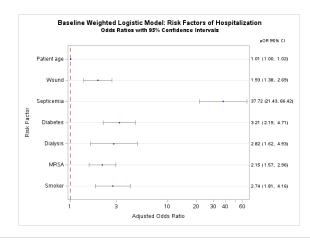
Type 3 Analysis of Effects					
Effect	F Value	Num DF	Den DF	Pr > F <.0001	
DIABETES	35.62	- 1	1445		
BSI	158.48	1	1445	<.0001	
WOUND	15.02	1	1445	0.0001	
newage	9.69	1	1445	0.0019	

Standard					
Parameter		Estimate	Error	t Value	Pr > t
Intercept		-2.3264	0.1864	-12.48	<.000
SMOKER	Yes	1.0095	0.2126	4.75	<.000
mrsafinal	MRSA	0.7667	0.1620	4.73	<.000
kidney	Yes	1.0379	0.2838	3.66	0.0003
DIABETES	Yes	1.1657	0.1953	5.97	<.000
BSI	Yes	3.6303	0.2884	12.59	<.000
WOUND	Yes	0.6579	0.1697	3.88	0.000
newage		0.0108	0.00346	3.11	0.0019

Odds Ratio Estimates			
Effect	Point Estimate	95% Confider	nce Limits
SMOKER Yes vs No	2.744	1.808	4.16
mrsafinal MRSA vs MSSA	2.153	1.567	2.95
kidney Yes vs No	2.823	1.618	4.92
DIABETES Yes vs No	3.208	2.187	4.70
BSI Yes vs No	37.725	21.427	66.42
WOUND Yes vs No	1.931	1.384	2.69
newage	1.011	1.004	1.01

Association of Predicted	Probabilities a	and Observed R	esponses
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	c	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



Weighted Logistic Stratified: Community-onset model

The SURVEYLOGISTIC Procedure

Model Infor	mation
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	Dograpo of Erondom (DE)		

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		
Class	Value	Design Variables
mrsafinal	MRSA	1
	MSSA	0
kidney	No	0
	Yes	1
DIABETES	No	0
	Yes	1
SMOKER	No	0
	Yes	1
BSI	No	0
	Yes	1
WOUND	No	0
	Yes	1

Stratum Information		
Stratum Index	INVASIVE	N Obs
1	No	679

Stratum Information		
Stratum Index	INVASIVE	N Obs
2	Yes	167

Model Convergence Status

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

Model Fit Statistics				
Criterion	Intercept Only	Intercept and Covariates		
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

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

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	<.000
SMOKER	Yes	1.2164	0.2678	4.54	<.000
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	<.000
BSI	Yes	4.5960	0.4142	11.10	<.000
WOUND	Yes	0.6934	0.2410	2.88	0.004
newage		0.00406	0.00504	0.81	0.420

Odds Ratio Estimates				
Effect	Point Estimate	stimate 95% Confide		
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	

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	c	0.837		

Weighted Logistic Stratified: Hospital-associated model

e SURVEYLOGISTIC Procedure

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	

Model Information		
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 Waights Head	1456

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

Probability modeled is hosp='Yes'.

Clas	s Level I	nformation
Class	Value	Design Variables
mrsafinal	MRSA	1
	MSSA	0
kidney	No	0
	Yes	1
DIABETES	No	0
	Yes	1
SMOKER	No	0
	Yes	1
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

	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	40.62	7	593	<.0001
Wald	10.58	7	593	<.0001

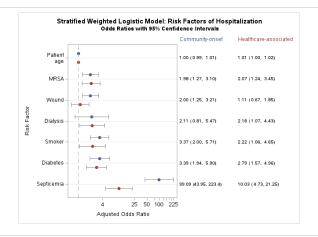
Type 3 Analysis of Effects					
Effect	F Value	Num DF	Den DF	Pr > F	
SMOKER	4.50	1	599	0.0342	
mrsafinal	7.76	- 1	599	0.0055	
kidney	4.68	1	599	0.0309	
DIABETES	12.32	1	599	0.0005	
BSI	36.32	1	599	<.0001	
WOUND	0.17	1	599	0.6819	
newage	1.27	1	599	0.2608	

			Standard		
Parameter		Estimate	Error	t Value	Pr > t
Intercept		-0.9211	0.3420	-2.69	0.007
SMOKER	Yes	0.7982	0.3761	2.12	0.034
mrsafinal	MRSA	0.7257	0.2606	2.79	0.005

Analysis of Maximum Likelihood Estimates					
Parameter		Estimate	Standard Error	t Value	Pr > t
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 deg	rees of freed	om for the t	tests is 5	99.

Odds Ratio Estimates				
Effect	Point Estimate	95% Confid	ence 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	

Association of Predicted F	Probabilities	and Observed R	esponses
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	c	0.815



Weighted Logistic Random-Intercept Model Conditional on Hospital Cluster

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 GA056 GA069 GA076 GA071 GA308 GAMDO OSODO

	Class Level Information		
Class	Levels	Values	
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

Ordered Value	hosp	Tota Frequency
1	No	735
2	Yes	712

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 Info	ormation
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

		Itera	tion 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.

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

	Covari	ance Param	eter Estimal	tes	
Cov Parm	Subject	Estimate	Standard Error	Z Value	Pr > Z
Intercept	TXHOSP	3.3699	2.2234	1.52	0.0648

				Solutions	for Fixed E	ffects					
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t
Intercept							-2.2875	0.5930	34	-3.86	0.000
SMOKER				Yes			0.5006	0.3209	18	1.56	0.136
SMOKER				No			0				
mrsafinal	MRSA						0.01493	0.2164	24	0.07	0.945
mrsafinal	MSSA						0				
kidney		Yes					1.2379	0.3360	22	3.68	0.001
kidney		No					0				
DIABETES			Yes				0.7665	0.2814	22	2.72	0.012
DIABETES			No				0				
WOUND					Yes		0.4193	0.2014	21	2.08	0.049
WOUND					No		0				
BSI						Yes	2.7266	0.3766	24	7.24	<.000
BSI						No	0				
newage							0.02178	0.005370	4321	4.06	<.000

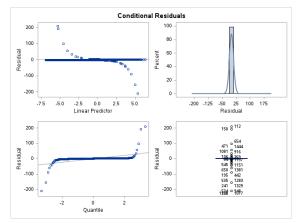
	Odds Ratio Estimates																
mrsafinal	safinal kidney DIABETES SMOKER WOUND BSI newage _mrsafinal _kidney _DIABETES _SMOKER _WOUND _BSI _newage Estimate DF 95% Confidence Limits																
			Yes			44.242				No			44.242	1.650	18	0.841	3.238
MRSA						44.242	MSSA						44.242	1.015	24	0.649	1.587
	Yes					44.242		No					44.242	3.448	22	1.718	6.923
		Yes				44.242			No				44.242	2.152	22	1.201	3.858
				Yes		44.242					No		44.242	1.521	21	1.000	2.312
					Yes	44.242						No	44.242	15.281	24	7.024	33.242
						45.242							44.242	1.022	4321	1.011	1.03

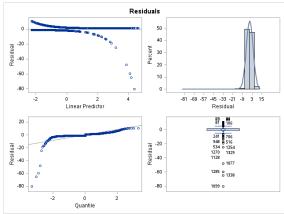
Type III Tests of Fixed Effects								
Effect	Num DF	Den DF	Chi-Square	F Value	Pr > ChiSq	Pr > F		
SMOKER	1	18	2.43	2.43	0.1188	0.1362		
mrsafinal	1	24	0.00	0.00	0.9450	0.9456		
kidney	1	22	13.57	13.57	0.0002	0.0013		
DIABETES	1	22	7.42	7.42	0.0065	0.0124		
WOUND	1	21	4.33	4.33	0.0374	0.0498		
BSI	1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

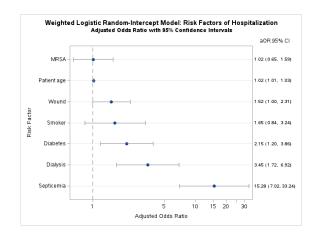
			Solution for Ra	ndom E	ffects				
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept	TXHOSP GA002	1.0334	0.8120	4355	1.27	0.2032	0.05	-0.5585	2.6253
Intercept	TXHOSP GA003	0.1329	0.6562	4355	0.20	0.8395	0.05	-1.1536	1.4194
Intercept	TXHOSP GA004	1.5172	0.5979	4355	2.54	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
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

			Solution for Ra	ndom E	ffects						
Effect	Subject	Subject Estimate Std Err Pred DF t Value Pr > t Alpha Lower Upper									
Intercept	TXHOSP OSODC	-1.8421	1.4661	4355	-1.26	0.2090	0.05	-4.7164	1.0322		

								Emp	irical Corre	lation Ma	trix for Fix	ed Effects	8								
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Row	Col1	Col2	Col3	Col4	Col5	Col6	Col7	Col8	Col9	Col10	Col11	Col12	Col13	Col14
Intercept							- 1	1.0000	-0.08803		-0.2056		-0.06619		-0.04415		-0.1236		-0.07947		-0.1829
SMOKER				Yes			2	-0.08803	1.0000		0.08159		0.03500		0.2710		0.3643		0.1116		-0.07769
SMOKER				No			3			1.0000											
mrsafinal	MRSA						4	-0.2056	0.08159		1.0000		0.1599		0.1505		0.2547		0.1467		-0.3066
mrsafinal	MSSA						5					1.0000									
kidney		Yes					6	-0.06619	0.03500		0.1599		1.0000		-0.2076		-0.2413		0.02663		-0.3054
kidney		No					7							1.0000							
DIABETES			Yes				8	-0.04415	0.2710		0.1505		-0.2076		1.0000		0.07682		-0.05413		-0.1068
DIABETES			No				9									1.0000					
WOUND					Yes		10	-0.1236	0.3643		0.2547		-0.2413		0.07682		1.0000		0.2771		0.03849
WOUND					No		- 11											1.0000			
BSI						Yes	12	-0.07947	0.1116		0.1467		0.02663		-0.05413		0.2771		1.0000		-0.04147
BSI						No	13													1.0000	
newage							14	-0.1829	-0.07769		-0.3066		-0.3054		-0.1068		0.03849		-0.04147		1.0000







Weighted Logistic Random-Intercept Stratified Model: Community-Onset

The GLIMMIX Procedu

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

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

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-Newtor
Parameters in Optimization	9
Lower Boundaries	1
Upper Boundaries	0
Fixed Effects	Not Profiled

Optimization	Information
Starting From	GLM estimates
Quadrature Points	1

Iteration History									
Iteration	Restarts	Evaluations	Objective Function	Change	Max Gradien				
0	0	4	1501.0917107		2753.62				
- 1	0	6	1480.828851	20.26285972	79.9683				
2	0	2	1446.3052458	34.52360519	25.974				
3	0	2	1438.6102748	7.69497099	25.2484				
4	0	4	1434.7206514	3.88962333	47.5742				
5	0	4	1431.4966568	3.22399465	12.6080				
6	0	3	1431.0656799	0.43097691	13.7122				
7	0	4	1430.0803129	0.98536704	5.96292				
8	0	3	1429.733555	0.34675782	14.8066				
9	0	3	1429.6672263	0.06632873	2.60606				
10	0	3	1429.6524119	0.01481443	1.81574				
11	0	2	1429.6338936	0.01851827	2.52729				
12	0	3	1429.6301493	0.00374429	0.27673				
13	0	3	1429.6300976	0.00005169	0.18416				
14	0	3	1429.6300848	0.00001285	0.01690				

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	144

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

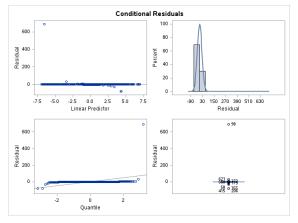
				Solutions	for Fixed E	ffects					
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t
Intercept							-2.5290	0.6961	29	-3.63	0.001
SMOKER				Yes			0.6718	0.4220	17	1.59	0.129
SMOKER				No			0				
mrsafinal	MRSA						-0.1810	0.2851	22	-0.64	0.532
mrsafinal	MSSA						0				
kidney		Yes					1.6130	0.6778	10	2.38	0.038
kidney		No					0				
DIABETES			Yes				1.1581	0.4643	17	2.49	0.023
DIABETES			No				0				
WOUND					Yes		0.5562	0.3572	16	1.56	0.139
WOUND					No		0				
BSI						Yes	4.0945	0.7998	19	5.12	<.000
BSI						No	0				
newage							0.02049	0.007379	2870	2.78	0.005

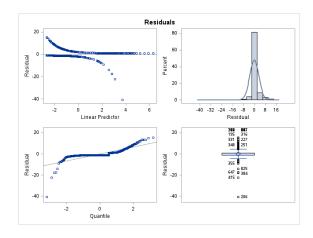
	Odds Ratio Estimates																
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BSI	_newage	Estimate	DF	95% Confid	ence Limits
			Yes			39.545				No			39.545	1.958	17	0.804	4.768
MRSA						39.545	MSSA						39.545	0.834	22	0.462	1.507
	Yes					39.545		No					39.545	5.018	10	1.108	22.721
		Yes				39.545			No				39.545	3.184	17	1.196	8.480
				Yes		39.545					No		39.545	1.744	16	0.818	3.719
					Yes	39.545						No	39.545	60.007	19	11.252	320.033
						40.545							39.545	1.021	2870	1.006	1.036
Effects of	continuou	s variables ar	re assessed	as one unit	offset	ts from the	mean. The A	T subontio	n modifies the	reference val	ue and the L	INIT sul	hontion mor	diffies the of	fsets.		

Type III Tests of Fixed Effects										
Effect	Num DF	Den DF	F Value	Pr>						
CMOKED	- 4	17	2.52	0.120						

Type III Tests of Fixed Effects										
Effect	Num DF	Den DF	F Value	Pr > F						
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	Upp	
Intercept	TXHOSP GA002	-1.3852	1.0465	2899	-1.32	0.1857	0.05	-3.4372	0.66	
Intercept	TXHOSP GA003	-0.6542	0.8484	2899	-0.77	0.4407	0.05	-2.3177	1.00	
Intercept	TXHOSP GA004	0.7617	0.6833	2899	1.11	0.2651	0.05	-0.5781	2.10	
Intercept	TXHOSP GA006	-1.3408	1.4341	2899	-0.93	0.3499	0.05	-4.1527	1.47	
Intercept	TXHOSP GA008	-1.6991	0.9333	2899	-1.82	0.0688	0.05	-3.5291	0.13	
Intercept	TXHOSP GA009	1.3989	0.8269	2899	1.69	0.0908	0.05	-0.2224	3.02	
Intercept	TXHOSP GA010	1.2832	0.6953	2899	1.85	0.0651	0.05	-0.08013	2.64	
Intercept	TXHOSP GA011	1.2810	0.6697	2899	1.91	0.0559	0.05	-0.03207	2.59	
Intercept	TXHOSP GA013	2.4816	0.8876	2899	2.80	0.0052	0.05	0.7413	4.22	
Intercept	TXHOSP GA015	2.3732	0.8225	2899	2.89	0.0039	0.05	0.7605	3.98	
Intercept	TXHOSP GA016	0.5821	1.6877	2899	0.34	0.7302	0.05	-2.7270	3.89	
Intercept	TXHOSP GA018	1.0116	0.8259	2899	1.22	0.2207	0.05	-0.6077	2.63	
Intercept	TXHOSP GA020	0.9734	0.6947	2899	1.40	0.1613	0.05	-0.3887	2.33	
Intercept	TXHOSP GA021	0.6896	0.7048	2899	0.98	0.3279	0.05	-0.6923	2.07	
Intercept	TXHOSP GA024	-0.09621	0.7157	2899	-0.13	0.8931	0.05	-1.4995	1.3	
Intercept	TXHOSP GA026	1.4177	0.7398	2899	1.92	0.0554	0.05	-0.03294	2.8	
Intercept	TXHOSP GA027	1.7194	0.7319	2899	2.35	0.0189	0.05	0.2842	3.1	
Intercept	TXHOSP GA030	-1.7086	1.3302	2899	-1.28	0.1991	0.05	-4.3168	0.8	
Intercept	TXHOSP GA032	1.5642	0.7456	2899	2.10	0.0360	0.05	0.1022	3.0	
Intercept	TXHOSP GA034	-1.4768	0.8780	2899	-1.68	0.0927	0.05	-3.1983	0.2	
Intercept	TXHOSP GA040	-4.0165	1.0574	2899	-3.80	0.0001	0.05	-6.0899	-1.9	
Intercept	TXHOSP GA046	2.8069	1.3843	2899	2.03	0.0427	0.05	0.09259	5.5	
Intercept	TXHOSP GA048	-0.7364	1.5905	2899	-0.46	0.6434	0.05	-3.8551	2.3	
Intercept	TXHOSP GA050	-2.4485	1.6212	2899	-1.51	0.1311	0.05	-5.6272	0.7	
Intercept	TXHOSP GA059	0.8361	0.7259	2899	1.15	0.2495	0.05	-0.5872	2.2	
Intercept	TXHOSP GA066	-0.7882	1.5712	2899	-0.50	0.6160	0.05	-3.8690	2.29	
Intercept	TXHOSP GA069	0.3367	1.8053	2899	0.19	0.8520	0.05	-3.2030	3.8	
Intercept	TXHOSP GA070	1.3729	0.8034	2899	1.71	0.0876	0.05	-0.2024	2.9	
Intercept	TXHOSP GA071	-1.0502	0.8703	2899	-1.21	0.2276	0.05	-2.7566	0.6	
Intercept	TXHOSP GAMDO	-4 2320	0.7083	2899	-5.97	< 0001	0.05	-5 6209	-28	





Weighted Logistic Random-Intercept Stratified Model: Hospital-Associated

The GLIMMIX Procedure

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=					

		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	
Total Frequency	hosp	Ordered Value
145	No	1
456	Yes	2
456 obability that hosp='Yes'.		2

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						

Optimization Information					
Fixed Effects	Not Profiled				
Starting From	GLM estimates				
Quadrature Points	5				

		Iterat	ion 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					
HOIC (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				

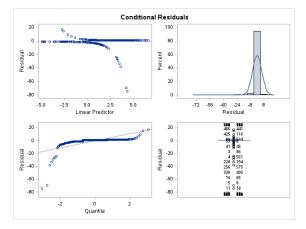
Covariance Parameter Estimates										
Cov Parm	Subject	Estimate	Standard Error	Z Value	Pr > Z					
Intercent	TXHOSP	4 0459	3 0019	1.35	0.0889					

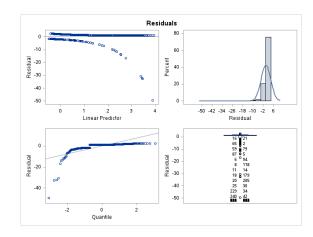
				Solutions	for Fixed E	ffects					
Effect	mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	Estimate	Standard Error	DF	t Value	Pr > t
Intercept							-0.4119	0.7588	30	-0.54	0.5912
SMOKER				Yes			0.4916	0.7261	16	0.68	0.5081
SMOKER				No			0				
mrsafinal	MRSA						0.2375	0.4195	22	0.57	0.5770
mrsafinal	MSSA						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

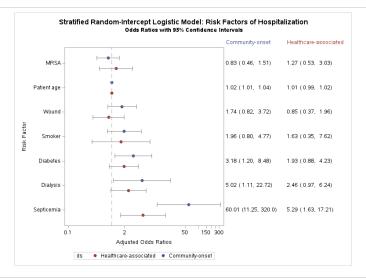
Odds Ratio Estimates																	
mrsafinal	kidney	DIABETES	SMOKER	WOUND	BSI	newage	_mrsafinal	_kidney	_DIABETES	_SMOKER	_WOUND	_BSI	_newage	Estimate	DF	95% Confidence Limits	
			Yes			53.62				No			53.62	1.635	16	0.351	7.620
MRSA						53.62	MSSA						53.62	1.268	22	0.531	3.027
	Yes					53.62		No					53.62	2.459	19	0.969	6.245
		Yes				53.62			No				53.62	1.934	19	0.884	4.232
				Yes		53.62					No		53.62	0.852	19	0.370	1.960
					Yes	53.62						No	53.62	5.290	21	1.626	17.212
						54.62							53.62	1.005	1418	0.990	1.021
Effects of o	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							

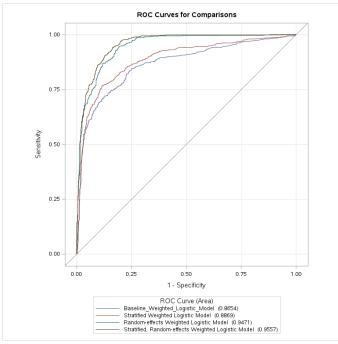
Solution for Random Effects											
Effect	Subject	Estimate	Std Err Pred	DF	t Value	Pr > t	Alpha	Lower	Upper		
Intercept	TXHOSP GA002	2.6027	1.3808	1448	1.88	0.0596	0.05	-0.1059	5.3113		
Intercept	TXHOSP GA003	0.5977	0.8245	1448	0.72	0.4686	0.05	-1.0196	2.2150		
Intercept	TXHOSP GA004	1.8489	0.7062	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		
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			Tau-a					
ROC Model	Area	Area Error Confidence Limits			Somers' D		Gamma				
Baseline_Weighted_Logistic_Model		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				

