The FREQ Procedure

Clinic

CLINIC	Frequency	Percent	Cumulative Frequency	Cumulative Percent
CPMC	12	9.23	12	9.23
SU	14	10.77	26	20.00
TTU	16	12.31	42	32.31
TU	32	24.62	74	56.92
UMI	3	2.31	77	59.23
UMS	40	30.77	117	90.00
WFU	13	10.00	130	100.00

Table of TX by primary

TX(Treatment group)
primary(Primary outcome: 2+ visits with FU_GCSI<=50%% BL_GCSI)

Frequency Percent Row Pct Col Pct	0	Total
Nor	65 50.00 100.00 50.00	65 50.00
Plb	65 50.00 100.00 50.00	65 50.00
Total	130 100.00	130 100.00

The MEANS Procedure

Variable	Label	N
btot	BL GCSI total score	130
bsub1	BL GCSI nausea subscore	130
bsub2	BL GCSI fullness subscore	130
bsub3	BL GCSI bloating subscore	130
bupain	BL Upper abdominal pain score	130
blpain	BL Lower abdominal pain score	130
bregurg	BL GERD subscore	130
bconstipa	BL consitpation score	130
bdiarrhea	BL diarrhea score	130
bpcs	BL SF-36 Physical Component Summary	130
bmcs	BL SF-36 Mental Component Summary	130
bbdi	BL Beck Depression Inventory	130
bsever	BL Brief Pain Inventory - severity score	130
binter	BL Brief Pain Inventory - interference score	130
bstate	BL State Anxiety Inventory score	130
btrait	BL Trait Anxiety Inventory score	130
bphq15	BL PHQ-15 score	130
bbmi	BL BMI - kg/m**2	130
bensure	BL Satiety volume - mL Ensure	130
bblbrad	BL Baseline power in bradygastria region - %	104
bblnorm	BL Baseline power in normal region - %	104
bbltach	BL Baseline power in tachygastria region - %	104
bblduod	BL Baseline power in duodenal region - %	104
bpstbrad	BL 0-30 min power in bradygastria region - %	104
bpstnorm	BL 0-30 min power in normal region - %	104
bpsttach	BL 0-30 min power in tachygastria region - %	104
bpstduod	BL 0-30 min power in duodenal region - %	104
tot	F15 GCSI total score	118
sub1	F15 GCSI nausea subscore	118
sub2	F15 GCSI fullness subscore	118
sub3	F15 GCSI bloating subscore	118
upain	F15 Upper abdominal pain score	118
lpain	F15 Lower abdominal pain score	118
regurg	F15 GERD subscore	118
constipa	F15 consitpation score	118
diarrhea	F15 diarrhea score	118
pcs	BL SF-36 Physical Component Summary	117
mcs	BL SF-36 Mental Component Summary	117
bdi	F15 Beck Depression Inventory	118
sever	F15 Brief Pain Inventory - severity score	117
inter	F15 Brief Pain Inventory - interference score	116
state	F15 State Anxiety Inventory score	118
trait	F15 Trait Anxiety Inventory score	117
phq15	F15 PHQ-15 score	118
bmi	F12 BMI - kg/m**2	114
ensure	F12 Satiety volume - mL Ensure	104
fblbrad	F12 Baseline power in bradygastria region - %	84

The MEANS Procedure

Variable	Label	N
fblnorm	F12 Baseline power in normal region - %	84
fbltach	F12 Baseline power in tachygastria region - %	84
fblduod	F12 Baseline power in duodenal region - %	84
fpstbrad	F12 0-30 min power in bradygastria region - %	85
fpstnorm	F12 0-30 min power in normal region - %	85
fpsttach	F12 0-30 min power in tachygastria region - %	85
fpstduod	F12 0-30 min power in duodenal region - %	85
ctot	F15-BL GCSI total score	118
csub1	F15-BL GCSI nausea subscore	118
csub2	F15-BL GCSI fullness subscore	118
csub3	F15-BL GCSI bloating subscore	118
cupain	F15-BL Upper abdominal pain score	118
clpain	F15-BL Lower abdominal pain score	118
cregurg	F15-BL GERD subscore	118
cconstipa	F15-BL consitpation score	118
cdiarrhea	F15-BL diarrhea score	118
cpcs	F15-BL SF-36 Physical Componet Summary	117
cmcs	F15-BL SF-36 Mental Componet Summary	117
cbdi	F15-BL Beck Depression Inventory	118
csever	F15-BL Brief Pain Inventory - severity score	117
cinter	F15-BL Brief Pain Inventory - interference score	116
cstate	F15-BL State Anxiety Inventory score	118
ctrait	F15-BL Trait Anxiety Inventory score	117
cphq15	F15-BL PHQ-15 score	118
cbmi	F12-BL BMI - kg/m**2	114
censure	F12-BL Satiety volume - mL Ensure	104
cblbrad	F12-BL Baseline power in bradygastria region - %	72
cblnorm	F12-BL Baseline power in normal region - %	72
cbltach	F12-BL Baseline power in tachygastria region - %	72
cblduod	F12-BL Baseline power in duodenal region - %	72
cpstbrad	F12-BL 0-30 min power in bradygastria region - %	73
cpstnorm	F12-BL 0-30 min power in normal region - %	73
cpsttach	F12-BL 0-30 min power in tachygastria region - %	73
cpstduod	F12-BL 0-30 min power in duodenal region - %	73

Variable	Label	Mean
btot	BL GCSI total score	30.6076923
bsub1	BL GCSI nausea subscore	8.1923077
bsub2	BL GCSI fullness subscore	15.2769231
bsub3	BL GCSI bloating subscore	7.1384615
bupain	BL Upper abdominal pain score	6.6230769
blpain	BL Lower abdominal pain score	4.5000000
bregurg	BL GERD subscore	16.6769231
bconstipa	BL consitpation score	2.5692308
bdiarrhea	BL diarrhea score	1.8923077

The MEANS Procedure

Variable	Label	Mean
bpcs	BL SF-36 Physical Component Summary	35.4314905
bmcs	BL SF-36 Mental Component Summary	40.1563803
bbdi	BL Beck Depression Inventory	17.4538462
bsever	BL Brief Pain Inventory - severity score	4.0365385
binter	BL Brief Pain Inventory - interference score	4.1241758
bstate	BL State Anxiety Inventory score	41.7384615
btrait	BL Trait Anxiety Inventory score	43.0538462
bphq15	BL PHQ-15 score	13.9923077
bbmi	BL BMI - kg/m**2	27.4743856
bensure	BL Satiety volume - mL Ensure	304.7153846
bblbrad	BL Baseline power in bradygastria region - %	46.1730769
bblnorm	BL Baseline power in normal region - %	19.8081731
bbltach	BL Baseline power in tachygastria region - %	23.3746154
bblduod	BL Baseline power in duodenal region - %	10.6439423
bpstbrad	BL 0-30 min power in bradygastria region - %	40.5543269
bpstnorm	BL 0-30 min power in normal region - %	23.4562500
bpsttach	BL 0-30 min power in tachygastria region - %	27.4599038
bpstduod	BL 0-30 min power in duodenal region - %	8.5291827
tot	F15 GCSI total score	22.1355932
sub1	F15 GCSI nausea subscore	5.3220339
sub2	F15 GCSI fullness subscore	11.0254237
sub3	F15 GCSI bloating subscore	5.7881356
upain	F15 Upper abdominal pain score	4.8474576
lpain	F15 Lower abdominal pain score	3.7796610
regurg	F15 GERD subscore	11.2372881
constipa	F15 consitpation score	2.1779661
diarrhea	F15 diarrhea score	1.3389831
pcs	BL SF-36 Physical Component Summary	38.5759599
mcs	BL SF-36 Mental Component Summary	42.1595185
bdi	F15 Beck Depression Inventory	13.6271186
sever	F15 Brief Pain Inventory - severity score	3.1474359
inter	F15 Brief Pain Inventory - interference score	3.3004926
state	F15 State Anxiety Inventory score	41.2796610
trait	F15 Trait Anxiety Inventory score	41.3504274
phq15	F15 PHQ-15 score	11.7542373
bmi	F12 BMI - kg/m**2	27.7084716
ensure	F12 Satiety volume - mL Ensure	311.5769231
fblbrad	F12 Baseline power in bradygastria region - %	50.6385714
fblnorm	F12 Baseline power in normal region - %	18.7732143
fbltach	F12 Baseline power in tachygastria region - %	21.2727381
fblduod	F12 Baseline power in duodenal region - %	9.3140476
fpstbrad	F12 0-30 min power in bradygastria region - %	40.7961765
fpstnorm	F12 0-30 min power in normal region - %	23.0592941
fpsttach	F12 0-30 min power in tachygastria region - %	27.3015882
fpstduod	F12 0-30 min power in duodenal region - %	8.8368824
ctot	F15-BL GCSI total score	-7.9830508
csub1	F15-BL GCSI nausea subscore	-2.6355932

The MEANS Procedure

Variable	Label	Mean
csub2	F15-BL GCSI fullness subscore	-4.1271186
csub3	F15-BL GCSI bloating subscore	-1.2203390
cupain	F15-BL Upper abdominal pain score	-1.7033898
clpain	F15-BL Lower abdominal pain score	-0.5508475
cregurg	F15-BL GERD subscore	-5.0084746
cconstipa	F15-BL consitpation score	-0.3389831
cdiarrhea	F15-BL diarrhea score	-0.5169492
cpcs	F15-BL SF-36 Physical Componet Summary	2.7175695
cmcs	F15-BL SF-36 Mental Componet Summary	1.3568980
cbdi	F15-BL Beck Depression Inventory	-2.8474576
csever	F15-BL Brief Pain Inventory - severity score	-0.8034188
cinter	F15-BL Brief Pain Inventory - interference score	-0.5886700
cstate	F15-BL State Anxiety Inventory score	0.1440678
ctrait	F15-BL Trait Anxiety Inventory score	-1.0341880
cphq15	F15-BL PHQ-15 score	-1.9152542
cbmi	F12-BL BMI - kg/m**2	0.2439664
censure	F12-BL Satiety volume - mL Ensure	3.8461538
cblbrad	F12-BL Baseline power in bradygastria region - %	2.0208333
cblnorm	F12-BL Baseline power in normal region - %	-0.8720833
cbltach	F12-BL Baseline power in tachygastria region - %	-0.8701389
cblduod	F12-BL Baseline power in duodenal region - %	-0.2798611
cpstbrad	F12-BL 0-30 min power in bradygastria region - %	-1.6354795
cpstnorm	F12-BL 0-30 min power in normal region - %	-0.1807534
cpsttach	F12-BL 0-30 min power in tachygastria region - %	0.7813699
cpstduod	F12-BL 0-30 min power in duodenal region - %	1.0283562

Variable	Label	Std Dev
btot	BL GCSI total score	6.2796536
bsub1	BL GCSI nausea subscore	3.9258558
bsub2	BL GCSI fullness subscore	3.7437605
bsub3	BL GCSI bloating subscore	2.7779694
bupain	BL Upper abdominal pain score	2.8509903
blpain	BL Lower abdominal pain score	3.2996359
bregurg	BL GERD subscore	9.9691115
bconstipa	BL consitpation score	1.8339539
bdiarrhea	BL diarrhea score	1.7575801
bpcs	BL SF-36 Physical Component Summary	9.9584810
bmcs	BL SF-36 Mental Component Summary	13.0253147
bbdi	BL Beck Depression Inventory	11.6566590
bsever	BL Brief Pain Inventory - severity score	2.5516205
binter	BL Brief Pain Inventory - interference score	3.1124263
bstate	BL State Anxiety Inventory score	12.4791633
btrait	BL Trait Anxiety Inventory score	12.3459886
bphq15	BL PHQ-15 score	4.8632400
bbmi	BL BMI - kg/m**2	6.3566077

The MEANS Procedure

Variable	Label	Std Dev
bensure	BL Satiety volume - mL Ensure	159.3631334
bblbrad	BL Baseline power in bradygastria region - %	19.2595078
bblnorm	BL Baseline power in normal region - %	12.7538082
bbltach	BL Baseline power in tachygastria region - %	10.2887666
bblduod	BL Baseline power in duodenal region - %	11.5408807
bpstbrad	BL 0-30 min power in bradygastria region - %	13.6545814
bpstnorm	BL 0-30 min power in normal region - %	12.6405057
bpsttach	BL 0-30 min power in tachygastria region - %	8.4917627
bpstduod	BL 0-30 min power in duodenal region - %	7.2965249
tot	F15 GCSI total score	10.2472971
sub1	F15 GCSI nausea subscore	4.0864198
sub2	F15 GCSI fullness subscore	5.3708009
sub3	F15 GCSI bloating subscore	3.2419570
upain	F15 Upper abdominal pain score	3.1747592
lpain	F15 Lower abdominal pain score	3.3769849
regurg	F15 GERD subscore	9.1368307
constipa	F15 consitpation score	1.8286951
diarrhea	F15 diarrhea score	1.6341460
pcs	BL SF-36 Physical Component Summary	11.3423689
mcs	BL SF-36 Mental Component Summary	13.6655123
bdi	F15 Beck Depression Inventory	11.7019953
sever	F15 Brief Pain Inventory - severity score	2.6695025
inter	F15 Brief Pain Inventory - interference score	3.3499463
state	F15 State Anxiety Inventory score	14.5248891
trait	F15 Trait Anxiety Inventory score	13.5190263
phq15	F15 PHQ-15 score	5.4833648
bmi	F12 BMI - kg/m**2	6.3521530
ensure	F12 Satiety volume - mL Ensure	169.2581009
fblbrad	F12 Baseline power in bradygastria region - %	19.9069951
fblnorm	F12 Baseline power in normal region - %	13.3844389
fbltach	F12 Baseline power in tachygastria region - %	10.7113610
fblduod	F12 Baseline power in duodenal region - %	10.2062162
fpstbrad	F12 0-30 min power in bradygastria region - %	14.5293176
fpstnorm	F12 0-30 min power in normal region - %	13.2611603
fpsttach	F12 0-30 min power in tachygastria region - %	10.1022680
fpstduod	F12 0-30 min power in duodenal region - %	9.5986796
ctot	F15-BL GCSI total score	10.0196244
csub1	F15-BL GCSI nausea subscore	3.9563111
csub2	F15-BL GCSI fullness subscore	5.3342092
csub3	F15-BL GCSI bloating subscore	3.0046562
cupain	F15-BL Upper abdominal pain score	3.2615589
clpain	F15-BL Lower abdominal pain score	3.0483215
cregurg	F15-BL GERD subscore	8.6661694
cconstipa	F15-BL consitpation score	1.5592039
cdiarrhea	F15-BL diarrhea score	1.6524163
cpcs	F15-BL SF-36 Physical Componet Summary	8.5516882
cmcs	F15-BL SF-36 Mental Componet Summary	10.3037960

The MEANS Procedure

Variable	Label	Std Dev
cbdi	F15-BL Beck Depression Inventory	8.0687507
csever	F15-BL Brief Pain Inventory - severity score	2.4567003
cinter	F15-BL Brief Pain Inventory - interference score	2.7655605
cstate	F15-BL State Anxiety Inventory score	11.0633571
ctrait	F15-BL Trait Anxiety Inventory score	8.6860288
cphq15	F15-BL PHQ-15 score	4.2073933
cbmi	F12-BL BMI - kg/m**2	1.2374807
censure	F12-BL Satiety volume - mL Ensure	120.3678641
cblbrad	F12-BL Baseline power in bradygastria region - %	23.9157451
cblnorm	F12-BL Baseline power in normal region - %	14.8631547
cbltach	F12-BL Baseline power in tachygastria region - %	13.0643047
cblduod	F12-BL Baseline power in duodenal region - %	12.8305489
cpstbrad	F12-BL 0-30 min power in bradygastria region - %	16.2094858
cpstnorm	F12-BL 0-30 min power in normal region - %	12.7151876
cpsttach	F12-BL 0-30 min power in tachygastria region - %	9.7264872
cpstduod	F12-BL 0-30 min power in duodenal region - %	10.7022138

Variable	Label	Minimum
btot	BL GCSI total score	21.0000000
bsub1	BL GCSI nausea subscore	1.0000000
bsub2	BL GCSI fullness subscore	3.0000000
bsub3	BL GCSI bloating subscore	0
bupain	BL Upper abdominal pain score	0
blpain	BL Lower abdominal pain score	0
bregurg	BL GERD subscore	0
bconstipa	BL consitpation score	0
bdiarrhea	BL diarrhea score	0
bpcs	BL SF-36 Physical Component Summary	10.4317639
bmcs	BL SF-36 Mental Component Summary	9.0928959
bbdi	BL Beck Depression Inventory	0
bsever	BL Brief Pain Inventory - severity score	0
binter	BL Brief Pain Inventory - interference score	0
bstate	BL State Anxiety Inventory score	20.0000000
btrait	BL Trait Anxiety Inventory score	20.0000000
bphq15	BL PHQ-15 score	2.0000000
bbmi	BL BMI - kg/m**2	16.6848256
bensure	BL Satiety volume - mL Ensure	40.0000000
bblbrad	BL Baseline power in bradygastria region - %	4.4100000
bblnorm	BL Baseline power in normal region - %	3.9200000
bbltach	BL Baseline power in tachygastria region - %	4.1800000
bblduod	BL Baseline power in duodenal region - %	0.3100000
bpstbrad	BL 0-30 min power in bradygastria region - %	8.9250000
bpstnorm	BL 0-30 min power in normal region - %	4.9000000
bpsttach	BL 0-30 min power in tachygastria region - %	8.8650000
bpstduod	BL 0-30 min power in duodenal region - %	1.5450000

The MEANS Procedure

Variable	Label	Minimum
tot	F15 GCSI total score	0
sub1	F15 GCSI nausea subscore	0
sub2	F15 GCSI fullness subscore	0
sub3	F15 GCSI bloating subscore	0
upain	F15 Upper abdominal pain score	0
lpain	F15 Lower abdominal pain score	0
regurg	F15 GERD subscore	0
constipa	F15 consitpation score	0
diarrhea	F15 diarrhea score	0
pcs	BL SF-36 Physical Component Summary	15.8441247
mcs	BL SF-36 Mental Component Summary	5.2890981
bdi	F15 Beck Depression Inventory	0
sever	F15 Brief Pain Inventory - severity score	0
inter	F15 Brief Pain Inventory - interference score	0
state	F15 State Anxiety Inventory score	20.0000000
trait	F15 Trait Anxiety Inventory score	20.0000000
phq15	F15 PHQ-15 score	0
bmi	F12 BMI - kg/m**2	16.1688001
ensure	F12 Satiety volume - mL Ensure	20.0000000
fblbrad	F12 Baseline power in bradygastria region - %	6.7000000
fblnorm	F12 Baseline power in normal region - %	3.1100000
fbltach	F12 Baseline power in tachygastria region - %	5.4700000
fblduod	F12 Baseline power in duodenal region - %	0.9200000
fpstbrad	F12 0-30 min power in bradygastria region - %	6.7150000
fpstnorm	F12 0-30 min power in normal region - %	2.8350000
fpsttach	F12 0-30 min power in tachygastria region - %	8.5650000
fpstduod	F12 0-30 min power in duodenal region - %	1.7100000
ctot	F15-BL GCSI total score	-37.0000000
csub1	F15-BL GCSI nausea subscore	-14.0000000
csub2	F15-BL GCSI fullness subscore	-17.0000000
csub3	F15-BL GCSI bloating subscore	-8.0000000
cupain	F15-BL Upper abdominal pain score	-10.0000000
clpain	F15-BL Lower abdominal pain score	-8.0000000
cregurg	F15-BL GERD subscore	-29.0000000
cconstipa	F15-BL consitpation score	-5.0000000
cdiarrhea	F15-BL diarrhea score	-5.0000000
cpcs	F15-BL SF-36 Physical Componet Summary	-16.0592375
cmcs	F15-BL SF-36 Mental Componet Summary	-27.2549223
cbdi	F15-BL Beck Depression Inventory	-25.0000000
csever	F15-BL Brief Pain Inventory - severity score	-7.0000000
cinter	F15-BL Brief Pain Inventory - interference score	-6.7142857
cstate	F15-BL State Anxiety Inventory score	-35.0000000
ctrait	F15-BL Trait Anxiety Inventory score	-32.0000000
cphq15	F15-BL PHQ-15 score	-12.0000000
cbmi	F12-BL BMI - kg/m**2	-4.1689164
censure	F12-BL Satiety volume - mL Ensure	-450.0000000
cblbrad	F12-BL Baseline power in bradygastria region - %	-53.5500000

The MEANS Procedure

Variable	Label	Minimum
cblnorm	F12-BL Baseline power in normal region - %	-48.6500000
cbltach	F12-BL Baseline power in tachygastria region - %	-30.2900000
cblduod	F12-BL Baseline power in duodenal region - %	-35.3900000
cpstbrad	F12-BL 0-30 min power in bradygastria region - %	-45.1500000
cpstnorm	F12-BL 0-30 min power in normal region - %	-38.0000000
cpsttach	F12-BL 0-30 min power in tachygastria region - %	-19.7850000
cpstduod	F12-BL 0-30 min power in duodenal region - %	-34.3450000

Variable	Label	Maximum
btot	BL GCSI total score	45.0000000
bsub1	BL GCSI nausea subscore	15.0000000
bsub2	BL GCSI fullness subscore	20.0000000
bsub3	BL GCSI bloating subscore	10.0000000
bupain	BL Upper abdominal pain score	10.0000000
blpain	BL Lower abdominal pain score	10.0000000
bregurg	BL GERD subscore	35.0000000
bconstipa	BL consitpation score	5.0000000
bdiarrhea	BL diarrhea score	5.0000000
bpcs	BL SF-36 Physical Component Summary	57.8009737
bmcs	BL SF-36 Mental Component Summary	67.1439641
bbdi	BL Beck Depression Inventory	57.0000000
bsever	BL Brief Pain Inventory - severity score	10.0000000
binter	BL Brief Pain Inventory - interference score	10.0000000
bstate	BL State Anxiety Inventory score	75.0000000
btrait	BL Trait Anxiety Inventory score	74.0000000
bphq15	BL PHQ-15 score	27.0000000
bbmi	BL BMI - kg/m**2	45.8898169
bensure	BL Satiety volume - mL Ensure	750.0000000
bblbrad	BL Baseline power in bradygastria region - %	87.6400000
bblnorm	BL Baseline power in normal region - %	73.2600000
bbltach	BL Baseline power in tachygastria region - %	52.7700000
bblduod	BL Baseline power in duodenal region - %	60.5300000
bpstbrad	BL 0-30 min power in bradygastria region - %	75.1350000
bpstnorm	BL 0-30 min power in normal region - %	70.5600000
bpsttach	BL 0-30 min power in tachygastria region - %	53.9050000
bpstduod	BL 0-30 min power in duodenal region - %	42.5650000
tot	F15 GCSI total score	43.0000000
sub1	F15 GCSI nausea subscore	15.0000000
sub2	F15 GCSI fullness subscore	20.0000000
sub3	F15 GCSI bloating subscore	10.0000000
upain	F15 Upper abdominal pain score	10.0000000
lpain	F15 Lower abdominal pain score	10.0000000
regurg	F15 GERD subscore	35.0000000
constipa	F15 consitpation score	5.0000000
diarrhea	F15 diarrhea score	5.0000000

The MEANS Procedure

Variable	Label	Maximum
pcs	BL SF-36 Physical Component Summary	59.0100528
mcs	BL SF-36 Mental Component Summary	63.4109682
bdi	F15 Beck Depression Inventory	51.0000000
sever	F15 Brief Pain Inventory - severity score	9.0000000
inter	F15 Brief Pain Inventory - interference score	10.0000000
state	F15 State Anxiety Inventory score	79.0000000
trait	F15 Trait Anxiety Inventory score	74.0000000
phq15	F15 PHQ-15 score	25.0000000
bmi	F12 BMI - kg/m**2	44.9461491
ensure	F12 Satiety volume - mL Ensure	900.0000000
fblbrad	F12 Baseline power in bradygastria region - %	89.8700000
fblnorm	F12 Baseline power in normal region - %	75.4500000
fbltach	F12 Baseline power in tachygastria region - %	50.6800000
fblduod	F12 Baseline power in duodenal region - %	50.4600000
fpstbrad	F12 0-30 min power in bradygastria region - %	76.0550000
fpstnorm	F12 0-30 min power in normal region - %	67.9650000
fpsttach	F12 0-30 min power in tachygastria region - %	54.1050000
fpstduod	F12 0-30 min power in duodenal region - %	76.0250000
ctot	F15-BL GCSI total score	17.0000000
csub1	F15-BL GCSI nausea subscore	7.0000000
csub2	F15-BL GCSI fullness subscore	11.0000000
csub3	F15-BL GCSI bloating subscore	6.0000000
cupain	F15-BL Upper abdominal pain score	5.0000000
clpain	F15-BL Lower abdominal pain score	8.0000000
cregurg	F15-BL GERD subscore	21.0000000
cconstipa	F15-BL consitpation score	3.0000000
cdiarrhea	F15-BL diarrhea score	5.0000000
cpcs	F15-BL SF-36 Physical Componet Summary	26.8792496
cmcs	F15-BL SF-36 Mental Componet Summary	32.3564838
cbdi	F15-BL Beck Depression Inventory	26.0000000
csever	F15-BL Brief Pain Inventory - severity score	6.0000000
cinter	F15-BL Brief Pain Inventory - interference score	9.0000000
cstate	F15-BL State Anxiety Inventory score	27.0000000
ctrait	F15-BL Trait Anxiety Inventory score	20.0000000
cphq15	F15-BL PHQ-15 score	7.0000000
cbmi	F12-BL BMI - kg/m**2	4.5090999
censure	F12-BL Satiety volume - mL Ensure	300.0000000
cblbrad	F12-BL Baseline power in bradygastria region - %	46.1200000
cblnorm	F12-BL Baseline power in normal region - %	40.2700000
cbltach	F12-BL Baseline power in tachygastria region - %	29.2500000
cblduod	F12-BL Baseline power in duodenal region - %	39.4100000
cpstbrad	F12-BL 0-30 min power in bradygastria region - %	35.4400000
cpstnorm	F12-BL 0-30 min power in normal region - %	43.5850000
cpsttach	F12-BL 0-30 min power in tachygastria region - %	30.9050000
cpstduod	F12-BL 0-30 min power in duodenal region - %	61.2250000

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	ctot F15-BL GCSI total score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	10874.9891	94.5651
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	10874.9891	94.5651
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-434.3234	
Full Log Likelihood		-434.3234	
AIC (smaller is better)		876.6468	
AICC (smaller is better)		877.0008	
BIC (smaller is better)		887.7295	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	5.6208	4.5349	-3.2674	14.5090	1.54
TX	Nor	1	-1.2778	1.7724	-4.7516	2.1961	0.52
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
btot		1	-0.4315	0.1465	-0.7188	-0.1443	8.67
Scale		1	9.6000	0.6249	8.4502	10.9064	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.2152
TX	Nor	0.4710
TX	Plb	.
btot		0.0032
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	csub1	F15-BL GCSI nausea subscore

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	1447.4588	12.5866
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	1447.4588	12.5866
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-315.3407	
Full Log Likelihood		-315.3407	
AIC (smaller is better)		638.6814	
AICC (smaller is better)		639.0353	
BIC (smaller is better)		649.7641	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.9394	0.7954	-0.6194	2.4983	1.40
TX	Nor	1	0.2775	0.6457	-0.9882	1.5431	0.18
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bsub1		1	-0.4658	0.0834	-0.6293	-0.3023	31.17
Scale		1	3.5024	0.2280	3.0829	3.9790	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.2375
TX	Nor	0.6674
TX	Plb	.
bsub1		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	csub2	F15-BL GCSI fullness subscore

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	2866.2542	24.9239
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	2866.2542	24.9239
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-355.6493	
Full Log Likelihood		-355.6493	
AIC (smaller is better)		719.2985	
AICC (smaller is better)		719.6525	
BIC (smaller is better)		730.3813	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	3.7564	1.8963	0.0397	7.4730	3.92
TX	Nor	1	-1.4440	0.9102	-3.2281	0.3400	2.52
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bsub2		1	-0.4750	0.1198	-0.7099	-0.2402	15.71
Scale		1	4.9285	0.3208	4.3382	5.5992	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0476
TX	Nor	0.1126
TX	Plb	.
bsub2		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	csub3	F15-BL GCSI bloating subscore

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	905.7846	7.8764
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	905.7846	7.8764
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-287.6836	
Full Log Likelihood		-287.6836	
AIC (smaller is better)		583.3673	
AICC (smaller is better)		583.7213	
BIC (smaller is better)		594.4500	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	1.6605	0.7288	0.2320	3.0890	5.19
TX	Nor	1	-0.0897	0.5108	-1.0909	0.9115	0.03
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bsub3		1	-0.4050	0.0916	-0.5845	-0.2254	19.55
Scale		1	2.7706	0.1803	2.4387	3.1476	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0227
TX	Nor	0.8607
TX	Plb	.
bsub3		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cupain	F15-BL Upper abdominal pain score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	965.7401	8.3977
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	965.7401	8.3977
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-291.4651	
Full Log Likelihood		-291.4651	
AIC (smaller is better)		590.9303	
AICC (smaller is better)		591.2843	
BIC (smaller is better)		602.0130	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	1.7430	0.6990	0.3729	3.1131	6.22
TX	Nor	1	0.1021	0.5274	-0.9316	1.1358	0.04
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bupain		1	-0.5335	0.0914	-0.7127	-0.3543	34.05
Scale		1	2.8608	0.1862	2.5181	3.2501	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0127
TX	Nor	0.8465
TX	Plb	.
bupain		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	clpain	F15-BL Lower abdominal pain score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	865.1407	7.5230
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	865.1407	7.5230
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-284.9750	
Full Log Likelihood		-284.9750	
AIC (smaller is better)		577.9500	
AICC (smaller is better)		578.3040	
BIC (smaller is better)		589.0327	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	1.4664	0.4722	0.5409	2.3920	9.64
TX	Nor	1	-0.5625	0.4992	-1.5409	0.4159	1.27
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
blpain		1	-0.4042	0.0752	-0.5515	-0.2569	28.92
Scale		1	2.7077	0.1763	2.3834	3.0762	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0019
TX	Nor	0.2598
TX	Plb	.
blpain		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cregurg F15-BL GERD subscore

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	6414.1527	55.7752
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	6414.1527	55.7752
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-403.1738	
Full Log Likelihood		-403.1738	
AIC (smaller is better)		814.3476	
AICC (smaller is better)		814.7016	
BIC (smaller is better)		825.4304	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	2.2679	1.5260	-0.7230	5.2588	2.21
TX	Nor	1	0.2559	1.3681	-2.4255	2.9372	0.03
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bregurg		1	-0.4554	0.0696	-0.5918	-0.3189	42.77
Scale		1	7.3727	0.4799	6.4896	8.3760	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.1372
TX	Nor	0.8516
TX	Plb	.
bregurg		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cconstipa F15-BL consitpation score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	227.1731	1.9754
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	227.1731	1.9754
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-206.0814	
Full Log Likelihood		-206.0814	
AIC (smaller is better)		420.1627	
AICC (smaller is better)		420.5167	
BIC (smaller is better)		431.2455	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.4482	0.2406	-0.0233	0.9197	3.47
TX	Nor	1	0.3315	0.2569	-0.1720	0.8350	1.67
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bconstipa		1	-0.3753	0.0695	-0.5116	-0.2390	29.11
Scale		1	1.3875	0.0903	1.2213	1.5763	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0625
TX	Nor	0.1969
TX	Plb	.
bconstipa		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cdiarrhea	F15-BL diarrhea score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	225.8895	1.9643
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	225.8895	1.9643
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-205.7471	
Full Log Likelihood		-205.7471	
AIC (smaller is better)		419.4941	
AICC (smaller is better)		419.8481	
BIC (smaller is better)		430.5769	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.3676	0.2306	-0.0844	0.8195	2.54
TX	Nor	1	0.1161	0.2565	-0.3867	0.6188	0.20
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bdiarrhea		1	-0.5063	0.0735	-0.6503	-0.3622	47.46
Scale		1	1.3836	0.0901	1.2179	1.5719	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.1110
TX	Nor	0.6510
TX	Plb	.
bdiarrhea		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	ccgpi F15-BL CGPI

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	177.7279	1.5455
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	177.7279	1.5455
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-191.5993	
Full Log Likelihood		-191.5993	
AIC (smaller is better)		391.1987	
AICC (smaller is better)		391.5526	
BIC (smaller is better)		402.2814	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.4687	0.1727	0.1303	0.8071	7.37
TX	Nor	1	0.3128	0.2263	-0.1308	0.7564	1.91
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bcgpi		1	-0.7058	0.1124	-0.9260	-0.4856	39.47
Scale		1	1.2273	0.0799	1.0803	1.3943	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0066
TX	Nor	0.1669
TX	Plb	.
bcgpi		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	ctotgsrs F15-BL GSRS

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	95.2700	0.8284
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	95.2700	0.8284
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-154.8105	
Full Log Likelihood		-154.8105	
AIC (smaller is better)		317.6210	
AICC (smaller is better)		317.9750	
BIC (smaller is better)		328.7038	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.6814	0.2892	0.1146	1.2481	5.55
TX	Nor	1	-0.0488	0.1660	-0.3741	0.2765	0.09
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
btotgsrs		1	-0.3378	0.0730	-0.4809	-0.1946	21.38
Scale		1	0.8985	0.0585	0.7909	1.0208	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0185
TX	Nor	0.7688
TX	Plb	.
btotgsrs		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cpcs F15-BL SF-36 Physical Componet Summary

Number of Observations Read	130
Number of Observations Used	117
Missing Values	13

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	114	7888.9322	69.2012
Scaled Deviance	114	117.0000	1.0263
Pearson Chi-Square	114	7888.9322	69.2012
Scaled Pearson X2	114	117.0000	1.0263
Log Likelihood		-412.3618	
Full Log Likelihood		-412.3618	
AIC (smaller is better)		832.7235	
AICC (smaller is better)		833.0807	
BIC (smaller is better)		843.7722	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	9.1281	3.0201	3.2088	15.0473	9.14
TX	Nor	1	1.9573	1.5225	-1.0268	4.9414	1.65
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bpcs		1	-0.2044	0.0781	-0.3576	-0.0513	6.84
Scale		1	8.2114	0.5368	7.2239	9.3339	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0025
TX	Nor	0.1986
TX	Plb	.
bpcs		0.0089
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cmcs F15-BL SF-36 Mental Componet Summary

Number of Observations Read	130
Number of Observations Used	117
Missing Values	13

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	114	11065.2137	97.0633
Scaled Deviance	114	117.0000	1.0263
Pearson Chi-Square	114	11065.2137	97.0633
Scaled Pearson X2	114	117.0000	1.0263
Log Likelihood		-432.1550	
Full Log Likelihood		-432.1550	
AIC (smaller is better)		872.3100	
AICC (smaller is better)		872.6671	
BIC (smaller is better)		883.3587	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	11.1592	3.0992	5.0848	17.2335	12.96
TX	Nor	1	1.3171	1.8048	-2.2202	4.8545	0.53
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bmcs		1	-0.2554	0.0709	-0.3945	-0.1164	12.96
Scale		1	9.7249	0.6357	8.5554	11.0543	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0003
TX	Nor	0.4655
TX	Plb	.
bmcs		0.0003
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cbdi F15-BL Beck Depression Inventory

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	6854.6475	59.6056
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	6854.6475	59.6056
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-407.0926	
Full Log Likelihood		-407.0926	
AIC (smaller is better)		822.1852	
AICC (smaller is better)		822.5392	
BIC (smaller is better)		833.2679	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.7380	1.4424	-2.0889	3.5650	0.26
TX	Nor	1	0.1909	1.4088	-2.5703	2.9522	0.02
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bbdi		1	-0.2231	0.0620	-0.3446	-0.1017	12.97
Scale		1	7.6217	0.4961	6.7088	8.6588	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.6089
TX	Nor	0.8922
TX	Plb	.
bbdi		0.0003
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	csever	F15-BL Brief Pain Inventory - severity score

Number of Observations Read	130
Number of Observations Used	117
Missing Values	13

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	114	547.0625	4.7988
Scaled Deviance	114	117.0000	1.0263
Pearson Chi-Square	114	547.0625	4.7988
Scaled Pearson X2	114	117.0000	1.0263
Log Likelihood		-256.2456	
Full Log Likelihood		-256.2456	
AIC (smaller is better)		520.4911	
AICC (smaller is better)		520.8483	
BIC (smaller is better)		531.5398	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	1.1770	0.4111	0.3713	1.9827	8.20
TX	Nor	1	-0.6665	0.4006	-1.4516	0.1186	2.77
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bsever		1	-0.4220	0.0768	-0.5724	-0.2715	30.20
Scale		1	2.1623	0.1414	1.9023	2.4579	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0042
TX	Nor	0.0961
TX	Plb	.
bsever		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cinter F15-BL Brief Pain Inventory - interference score

Number of Observations Read	130
Number of Observations Used	116
Missing Values	14

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	113	744.7878	6.5910
Scaled Deviance	113	116.0000	1.0265
Pearson Chi-Square	113	744.7878	6.5910
Scaled Pearson X2	113	116.0000	1.0265
Log Likelihood		-272.4484	
Full Log Likelihood		-272.4484	
AIC (smaller is better)		552.8968	
AICC (smaller is better)		553.2572	
BIC (smaller is better)		563.9112	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	1.0490	0.4355	0.1955	1.9026	5.80
TX	Nor	1	-0.8736	0.4712	-1.7971	0.0500	3.44
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
binter		1	-0.3146	0.0756	-0.4629	-0.1663	17.30
Scale		1	2.5339	0.1664	2.2279	2.8819	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0160
TX	Nor	0.0638
TX	Plb	.
binter		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cstate	F15-BL State Anxiety Inventory score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	13500.8780	117.3989
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	13500.8780	117.3989
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-447.0844	
Full Log Likelihood		-447.0844	
AIC (smaller is better)		902.1689	
AICC (smaller is better)		902.5229	
BIC (smaller is better)		913.2516	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	8.6340	3.5328	1.7098	15.5582	5.97
TX	Nor	1	0.5436	1.9723	-3.3220	4.4093	0.08
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bstate		1	-0.2127	0.0797	-0.3689	-0.0564	7.11
Scale		1	10.6965	0.6963	9.4152	12.1520	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0145
TX	Nor	0.7828
TX	Plb	.
bstate		0.0076
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	ctrait	F15-BL Trait Anxiety Inventory score

Number of Observations Read	130
Number of Observations Used	117
Missing Values	13

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	114	8386.7798	73.5682
Scaled Deviance	114	117.0000	1.0263
Pearson Chi-Square	114	8386.7798	73.5682
Scaled Pearson X2	114	117.0000	1.0263
Log Likelihood		-415.9417	
Full Log Likelihood		-415.9417	
AIC (smaller is better)		839.8835	
AICC (smaller is better)		840.2406	
BIC (smaller is better)		850.9322	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	4.0093	3.0024	-1.8753	9.8940	1.78
TX	Nor	1	1.2486	1.5706	-1.8297	4.3269	0.63
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
btrait		1	-0.1331	0.0648	-0.2602	-0.0060	4.21
Scale		1	8.4665	0.5535	7.4483	9.6239	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.1818
TX	Nor	0.4266
TX	Plb	.
btrait		0.0401
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cphq15 F15-BL PHQ-15 score

Number of Observations Read	130
Number of Observations Used	118
Missing Values	12

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	115	1898.0568	16.5048
Scaled Deviance	115	118.0000	1.0261
Pearson Chi-Square	115	1898.0568	16.5048
Scaled Pearson X2	115	118.0000	1.0261
Log Likelihood		-331.3309	
Full Log Likelihood		-331.3309	
AIC (smaller is better)		670.6618	
AICC (smaller is better)		671.0158	
BIC (smaller is better)		681.7446	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	1.7951	1.1924	-0.5420	4.1321	2.27
TX	Nor	1	-1.0251	0.7404	-2.4762	0.4259	1.92
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bphq15		1	-0.2358	0.0775	-0.3878	-0.0839	9.25
Scale		1	4.0106	0.2611	3.5302	4.5564	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.1322
TX	Nor	0.1662
TX	Plb	.
bphq15		0.0024
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	cbmi F12-BL BMI - kg/m**2

Number of Observations Read	130
Number of Observations Used	114
Missing Values	16

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	111	165.1728	1.4880
Scaled Deviance	111	114.0000	1.0270
Pearson Chi-Square	111	165.1728	1.4880
Scaled Pearson X2	111	114.0000	1.0270
Log Likelihood		-182.8943	
Full Log Likelihood		-182.8943	
AIC (smaller is better)		373.7885	
AICC (smaller is better)		374.1555	
BIC (smaller is better)		384.7333	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	0.5682	0.5309	-0.4724	1.6087	1.15
TX	Nor	1	0.4311	0.2279	-0.0155	0.8777	3.58
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bbmi		1	-0.0194	0.0179	-0.0545	0.0157	1.17
Scale		1	1.2037	0.0797	1.0572	1.3705	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.2845
TX	Nor	0.0585
TX	Plb	.
bbmi		0.2792
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3
Distribution	Normal
Link Function	Identity
Dependent Variable	censure F12-BL Satiety volume - mL Ensure

Number of Observations Read	130
Number of Observations Used	104
Missing Values	26

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	101	1344322.8377	13310.1271
Scaled Deviance	101	104.0000	1.0297
Pearson Chi-Square	101	1344322.8377	13310.1271
Scaled Pearson X2	101	104.0000	1.0297
Log Likelihood		-639.8541	
Full Log Likelihood		-639.8541	
AIC (smaller is better)		1287.7083	
AICC (smaller is better)		1288.1123	
BIC (smaller is better)		1298.2858	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	71.4658	25.9901	20.5261	122.4055	7.56
TX	Nor	1	8.2897	22.3398	-35.4955	52.0749	0.14
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bensure		1	-0.2324	0.0690	-0.3676	-0.0973	11.36
Scale		1	113.6934	7.8832	99.2464	130.2433	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0060
TX	Nor	0.7106
TX	Plb	.
bensure		0.0008
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cblbrad	F12-BL Baseline power in bradygastria region - %

Number of Observations Read	130
Number of Observations Used	72
Missing Values	58

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	69	25615.8800	371.2446
Scaled Deviance	69	72.0000	1.0435
Pearson Chi-Square	69	25615.8800	371.2446
Scaled Pearson X2	69	72.0000	1.0435
Log Likelihood		-313.6384	
Full Log Likelihood		-313.6384	
AIC (smaller is better)		635.2769	
AICC (smaller is better)		635.8739	
BIC (smaller is better)		644.3835	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	36.3059	5.8297	24.8799	47.7319	38.78
TX	Nor	1	-2.4184	4.5365	-11.3098	6.4729	0.28
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bblbrad		1	-0.6936	0.1107	-0.9106	-0.4767	39.26
Scale		1	18.8620	1.5718	16.0197	22.2086	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		<.0001
TX	Nor	0.5940
TX	Plb	.
bblbrad		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cpstbrad	F12-BL 0-30 min power in bradygastria region - %

Number of Observations Read	130
Number of Observations Used	73
Missing Values	57

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	70	12301.5375	175.7362
Scaled Deviance	70	73.0000	1.0429
Pearson Chi-Square	70	12301.5375	175.7362
Scaled Pearson X2	70	73.0000	1.0429
Log Likelihood		-290.7187	
Full Log Likelihood		-290.7187	
AIC (smaller is better)		589.4375	
AICC (smaller is better)		590.0257	
BIC (smaller is better)		598.5993	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	27.4833	5.1314	17.4260	37.5407	28.69
TX	Nor	1	-1.8962	3.0567	-7.8872	4.0947	0.38
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bpstbrad		1	-0.6883	0.1099	-0.9037	-0.4730	39.25
Scale		1	12.9813	1.0743	11.0375	15.2674	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		<.0001
TX	Nor	0.5350
TX	Plb	.
bpstbrad		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cblnorm	F12-BL Baseline power in normal region - %

Number of Observations Read	130
Number of Observations Used	72
Missing Values	58

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	69	11548.6955	167.3724
Scaled Deviance	69	72.0000	1.0435
Pearson Chi-Square	69	11548.6955	167.3724
Scaled Pearson X2	69	72.0000	1.0435
Log Likelihood		-284.9594	
Full Log Likelihood		-284.9594	
AIC (smaller is better)		577.9188	
AICC (smaller is better)		578.5158	
BIC (smaller is better)		587.0255	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	9.0103	3.1735	2.7903	15.2303	8.06
TX	Nor	1	2.4391	2.9964	-3.4338	8.3119	0.66
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bblnorm		1	-0.5543	0.1110	-0.7719	-0.3366	24.92
Scale		1	12.6649	1.0554	10.7564	14.9119	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0045
TX	Nor	0.4156
TX	Plb	.
bblnorm		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cpstnorm	F12-BL 0-30 min power in normal region - %

Number of Observations Read	130
Number of Observations Used	73
Missing Values	57

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	70	8986.2962	128.3757
Scaled Deviance	70	73.0000	1.0429
Pearson Chi-Square	70	8986.2962	128.3757
Scaled Pearson X2	70	73.0000	1.0429
Log Likelihood		-279.2569	
Full Log Likelihood		-279.2569	
AIC (smaller is better)		566.5138	
AICC (smaller is better)		567.1020	
BIC (smaller is better)		575.6756	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	10.5389	2.8431	4.9666	16.1112	13.74
TX	Nor	1	-0.8891	2.6134	-6.0112	4.2331	0.12
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bpstnorm		1	-0.4281	0.0934	-0.6112	-0.2451	21.01
Scale		1	11.0950	0.9182	9.4337	13.0489	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0002
TX	Nor	0.7337
TX	Plb	.
bpstnorm		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cbltach	F12-BL Baseline power in tachygastria region - %

Number of Observations Read	130
Number of Observations Used	72
Missing Values	58

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	69	7434.2500	107.7428
Scaled Deviance	69	72.0000	1.0435
Pearson Chi-Square	69	7434.2500	107.7428
Scaled Pearson X2	69	72.0000	1.0435
Log Likelihood		-269.1023	
Full Log Likelihood		-269.1023	
AIC (smaller is better)		546.2046	
AICC (smaller is better)		546.8016	
BIC (smaller is better)		555.3113	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	15.7133	3.5795	8.6975	22.7290	19.27
TX	Nor	1	0.9593	2.5156	-3.9711	5.8898	0.15
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bbltach		1	-0.7754	0.1228	-1.0161	-0.5346	39.84
Scale		1	10.1614	0.8468	8.6302	11.9643	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		<.0001
TX	Nor	0.7029
TX	Plb	.
bbltach		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cpsttach	F12-BL 0-30 min power in tachygastria region - %

Number of Observations Read	130
Number of Observations Used	73
Missing Values	57

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	70	5905.9629	84.3709
Scaled Deviance	70	73.0000	1.0429
Pearson Chi-Square	70	5905.9629	84.3709
Scaled Pearson X2	70	73.0000	1.0429
Log Likelihood		-263.9364	
Full Log Likelihood		-263.9364	
AIC (smaller is better)		535.8729	
AICC (smaller is better)		536.4611	
BIC (smaller is better)		545.0347	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	10.8596	3.7583	3.4934	18.2257	8.35
TX	Nor	1	1.4173	2.1147	-2.7274	5.5619	0.45
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bpsttach		1	-0.4010	0.1242	-0.6444	-0.1576	10.43
Scale		1	8.9946	0.7444	7.6478	10.5786	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0039
TX	Nor	0.5027
TX	Plb	.
bpsttach		0.0012
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cblduod	F12-BL Baseline power in duodenal region - %

Number of Observations Read	130
Number of Observations Used	72
Missing Values	58

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	69	7322.9612	106.1299
Scaled Deviance	69	72.0000	1.0435
Pearson Chi-Square	69	7322.9612	106.1299
Scaled Pearson X2	69	72.0000	1.0435
Log Likelihood		-268.5593	
Full Log Likelihood		-268.5593	
AIC (smaller is better)		545.1186	
AICC (smaller is better)		545.7156	
BIC (smaller is better)		554.2253	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	7.4867	2.0800	3.4100	11.5634	12.96
TX	Nor	1	-1.3828	2.3867	-6.0606	3.2950	0.34
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bblduod		1	-0.6888	0.1053	-0.8952	-0.4824	42.79
Scale		1	10.0850	0.8404	8.5653	11.8744	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0003
TX	Nor	0.5623
TX	Plb	.
bblduod		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The GENMOD Procedure

Model Information

Data Set	WORK.TABLE3	
Distribution	Normal	
Link Function	Identity	
Dependent Variable	cpstduod	F12-BL 0-30 min power in duodenal region - %

Number of Observations Read	130
Number of Observations Used	73
Missing Values	57

Class Level Information

Class	Levels	Values
TX	2	Nor Plb

Criteria For Assessing Goodness Of Fit

Criterion	DF	Value	Value/DF
Deviance	70	6748.3872	96.4055
Scaled Deviance	70	73.0000	1.0429
Pearson Chi-Square	70	6748.3872	96.4055
Scaled Pearson X2	70	73.0000	1.0429
Log Likelihood		-268.8034	
Full Log Likelihood		-268.8034	
AIC (smaller is better)		545.6068	
AICC (smaller is better)		546.1950	
BIC (smaller is better)		554.7686	

Algorithm converged.

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		DF	Estimate	Standard Error	Wald 95% Confidence Limits		Wald Chi-Square
Intercept		1	5.9745	2.0886	1.8810	10.0681	8.18
TX	Nor	1	1.0318	2.2615	-3.4008	5.4643	0.21
TX	Plb	0	0.0000	0.0000	0.0000	0.0000	.
bpstduod		1	-0.6688	0.1663	-0.9948	-0.3429	16.18
Scale		1	9.6148	0.7957	8.1751	11.3080	

The GENMOD Procedure

Analysis Of Maximum Likelihood Parameter Estimates

Parameter		Pr > ChiSq
Intercept		0.0042
TX	Nor	0.6482
TX	Plb	.
bpstduod		<.0001
Scale		

NOTE: The scale parameter was estimated by maximum likelihood.

The CONTENTS Procedure

Data Set Name	LIMITED.TABLE3	Observations	130
Member Type	DATA	Variables	194
Engine	V9	Indexes	0
Created	Monday, October 13, 2014 01:01:57 PM	Observation Length	1816
Last Modified	Monday, October 13, 2014 01:01:57 PM	Deleted Observations	0
Protection		Compressed	NO
Data Set Type		Sorted	YES
Label			
Data Representation	WINDOWS_64		
Encoding	wlatin1 Western (Windows)		

Engine/Host Dependent Information

Data Set Page Size	16384
Number of Data Set Pages	16
First Data Page	2
Max Obs per Page	9
Obs in First Data Page	5
Number of Data Set Repairs	0
Filename	e:\gpcrc\norig\limited use dataset\table3.sas7bdat
Release Created	9.0301M0
Host Created	X64_7PRO

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
2	CLINIC	Char	4			Clinic
4	TRDATETIME	Char	22			trDatetime
3	TX	Char	3			Treatment group
52	anyquery	Char	1		$1.	
133	b15brad	Num	8			
145	b15duod	Num	8			
137	b15norm	Num	8			
141	b15tach	Num	8			
134	b30brad	Num	8			
146	b30duod	Num	8			
138	b30norm	Num	8			
142	b30tach	Num	8			
108	bbdi	Num	8			BL Beck Depression Inventory
70	bbigstom	Num	8			
132	bblbrad	Num	8			BL Baseline power in bradygastria region - %

The CONTENTS Procedure

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
144	bblduod	Num	8			BL Baseline power in duodenal region - %
136	bblnorm	Num	8			BL Baseline power in normal region - %
69	bbloat	Num	8			
140	bbltach	Num	8			BL Baseline power in tachygastria region - %
126	bbmi	Num	8			BL BMI - kg/m**2
148	bcbrad	Num	8			
151	bcduod	Num	8			
98	bcgpi	Num	8			BL CGPI
149	bcnorm	Num	8			
78	bconstipa	Num	8			BL consitpation score
150	bctach	Num	8			
109	bdi	Num	8			F15 Beck Depression Inventory
79	bdiarrhea	Num	8			BL diarrhea score
129	bensure	Num	8			BL Satiety volume - mL Ensure
67	bfull	Num	8			
88	bigstom	Num	8			
112	binter	Num	8			BL Brief Pain Inventory - interference score
87	bloat	Num	8			
68	blossapp	Num	8			
76	blpain	Num	8			BL Lower abdominal pain score
103	bmcs	Num	8			BL SF-36 Mental Component Summary
127	bmi	Num	8			F12 BMI - kg/m**2
62	bnausea	Num	8			
66	bnofinish	Num	8			
102	bpcs	Num	8			BL SF-36 Physical Component Summary
123	bphq15	Num	8			BL PHQ-15 score
135	bpstbrad	Num	8			BL 0-30 min power in bradygastria region - %
147	bpstduod	Num	8			BL 0-30 min power in duodenal region - %
139	bpstnorm	Num	8			BL 0-30 min power in normal region - %
143	bpsttach	Num	8			BL 0-30 min power in tachygastria region - %
77	bregurg	Num	8			BL GERD subscore
63	bretch	Num	8			
111	bsever	Num	8			BL Brief Pain Inventory - severity score
117	bstate	Num	8			BL State Anxiety Inventory score

The CONTENTS Procedure

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
65	bstomfull	Num	8			
71	bsub1	Num	8			BL GCSI nausea subscore
72	bsub2	Num	8			BL GCSI fullness subscore
73	bsub3	Num	8			BL GCSI bloating subscore
74	btot	Num	8			BL GCSI total score
100	btotgsrs	Num	8			BL GSRS
118	btrait	Num	8			BL Trait Anxiety Inventory score
75	bupain	Num	8			BL Upper abdominal pain score
64	bvomit	Num	8			
110	cbdi	Num	8			F15-BL Beck Depression Inventory
172	cblbrad	Num	8			F12-BL Baseline power in bradygastria region - %
178	cblduod	Num	8			F12-BL Baseline power in duodenal region - %
174	cblnorm	Num	8			F12-BL Baseline power in normal region - %
176	cbltach	Num	8			F12-BL Baseline power in tachygastria region - %
128	cbmi	Num	8			F12-BL BMI - kg/m**2
193	ccgpi	Num	8			F15-BL CGPI
191	cconstipa	Num	8			F15-BL consitpation score
192	cdiarrhea	Num	8			F15-BL diarrhea score
131	censure	Num	8			F12-BL Satiety volume - mL Ensure
99	cgpi	Num	8			F15 CGPI
116	cinter	Num	8			F15-BL Brief Pain Inventory - interference score
189	clpain	Num	8			F15-BL Lower abdominal pain score
107	cmcs	Num	8			F15-BL SF-36 Mental Componet Summary
96	constipa	Num	8			F15 consitpation score
106	cpcs	Num	8			F15-BL SF-36 Physical Componet Summary
125	cphq15	Num	8			F15-BL PHQ-15 score
173	cpstbrad	Num	8			F12-BL 0-30 min power in bradygastria region - %
179	cpstduod	Num	8			F12-BL 0-30 min power in duodenal region - %
175	cpstnorm	Num	8			F12-BL 0-30 min power in normal region - %
177	cpsttach	Num	8			F12-BL 0-30 min power in tachygastria region - %
190	cregurg	Num	8			F15-BL GERD subscore

The CONTENTS Procedure

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
115	csever	Num	8			F15-BL Brief Pain Inventory - severity score
121	cstate	Num	8			F15-BL State Anxiety Inventory score
185	csub1	Num	8			F15-BL GCSI nausea subscore
186	csub2	Num	8			F15-BL GCSI fullness subscore
187	csub3	Num	8			F15-BL GCSI bloating subscore
184	ctot	Num	8			F15-BL GCSI total score
194	ctotgsrs	Num	8			F15-BL GSRS
122	ctrait	Num	8			F15-BL Trait Anxiety Inventory score
188	cupain	Num	8			F15-BL Upper abdominal pain score
180	dcbrad	Num	8			
183	dcduod	Num	8			
181	dcnorm	Num	8			
182	dctach	Num	8			
97	diarrhea	Num	8			F15 diarrhea score
130	ensure	Num	8			F12 Satiety volume - mL Ensure
58	equ	Num	8			
153	f15brad	Num	8			
165	f15duod	Num	8			
157	f15norm	Num	8			
161	f15tach	Num	8			
154	f30brad	Num	8			
166	f30duod	Num	8			
158	f30norm	Num	8			
162	f30tach	Num	8			
152	fblbrad	Num	8			F12 Baseline power in bradygastria region - %
164	fblduod	Num	8			F12 Baseline power in duodenal region - %
156	fblnorm	Num	8			F12 Baseline power in normal region - %
160	fbltach	Num	8			F12 Baseline power in tachygastria region - %
168	fcbrad	Num	8			
171	fcduod	Num	8			
169	fcnorm	Num	8			
170	fctach	Num	8			
10	form	Char	4			6 Form/revision
8	formdate	Num	8	DATE7.		4 Date
155	fpstbrad	Num	8			F12 0-30 min power in bradygastria region - %
167	fpstduod	Num	8			F12 0-30 min power in duodenal region - %

The CONTENTS Procedure

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
159	fpstnorm	Num	8			F12 0-30 min power in normal region - %
163	fpsttach	Num	8			F12 0-30 min power in tachygastria region - %
85	full	Num	8			
14	gd108	Char	2			8 Nausea subscore
15	gd109	Char	2			9 Fullness/Satiety subscore
16	gd110	Char	2			10 Bloating subscore
17	gd111	Char	2			11 Total GSCI score
18	gd112	Char	1			12 s1 visit and GSCI score<20
21	gd114	Char	7			14 Date form reviewed
22	gd115	Char	1			15 Nausea
23	gd116	Char	1			16 Retching
24	gd117	Char	1			17 Vomiting
25	gd118	Char	1			18 Stomach fullness
26	gd119	Char	1			19 Unable to finish normal-size meal
27	gd120	Char	1			20 Feel full after meals
28	gd121	Char	1			21 Loss of appetite
29	gd122	Char	1			22 Bloating
30	gd123	Char	1			23 Stomach or belly visibly larger
31	gd124	Char	1			24 Upper abdominal pain
32	gd125	Char	1			25 Upper abdominal discomfort
33	gd126	Char	1			26 Lower abdominal pain
34	gd127	Char	1			27 Lower abdominal discomfort
35	gd128	Char	1			28 Heartburn during day
36	gd129	Char	1			29 Heartburn when lying down
37	gd130	Char	1			30 Chest discomfort during day
38	gd131	Char	1			31 Chest discomfort during night
39	gd132	Char	1			32 Regurgitation during day
40	gd133	Char	1			33 Regurgitation when lying down
41	gd134	Char	1			34 Bitter/sour/acid taste in mouth
42	gd135	Char	1			35 Constipation
43	gd136	Char	1			36 Diarrhea
44	gd137	Char	2			37 Predominant symptom
45	gd138	Char	7			38 Date form completed
19	gd113a	Char	3			13a Coordinator PIN
20	gd113b	Char	1			13b Coordinator signed (y=1)
1	id	Char	4			NORIG Limited use dataset ID number
114	inter	Num	8			F15 Brief Pain Inventory - interference score

The CONTENTS Procedure

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
53	key	Char	70			
86	lossapp	Num	8			
94	lpain	Num	8			F15 Lower abdominal pain score
105	mcs	Num	8			BL SF-36 Mental Component Summary
7	namecd	Char	3			3 Patient code
80	nausea	Num	8			
84	nofinish	Num	8			
60	nrec	Num	8			
12	oper	Char	8			
104	pcs	Num	8			BL SF-36 Physical Component Summary
124	phq15	Num	8			F15 PHQ-15 score
6	primary	Num	8			Primary outcome: 2+ visits with FU_GCSI<=50% BL_GCSI
46	pstudy	Char	10			
95	regurg	Num	8			F15 GERD subscore
81	retch	Num	8			
5	rzdate	Num	8	MMDDYY8.		
113	sever	Num	8			F15 Brief Pain Inventory - severity score
57	start	Num	8	DATETIME18.		
55	startk1	Num	8	DATETIME18.		
56	startk2	Num	8	DATETIME18.		
119	state	Num	8			F15 State Anxiety Inventory score
83	stomfull	Num	8			
11	study	Char	1			7 Study
89	sub1	Num	8			F15 GCSI nausea subscore
90	sub2	Num	8			F15 GCSI fullness subscore
91	sub3	Num	8			F15 GCSI bloating subscore
92	tot	Num	8			F15 GCSI total score
101	totgsrs	Num	8			F15 GSRS
61	totlen	Num	8			
120	trait	Num	8			F15 Trait Anxiety Inventory score
54	trantime	Num	8	DATETIME18.		
13	trantype	Char	3			
59	unequ	Num	8			
93	upain	Num	8			F15 Upper abdominal pain score
9	visit	Char	4			5 Visit ID code
82	vomit	Num	8			
47	zznote1	Char	175			
48	zznote2	Char	75			
49	zznote3	Char	75			
50	zznote4	Char	75			

The CONTENTS Procedure

Alphabetic List of Variables and Attributes

#	Variable	Type	Len	Format	Informat	Label
51	zznote5	Char	75			

Sort Information

Sortedby	id
Validated	YES
Character Set	ANSI