\*The data are from a prospective study on body fat accretion in a cohort of 162 girls from the MIT Growth and Development Study. The study examined changes in percent body fat before and after menarche. The data represent a subset of the study materials and should not be used to draw substantive conclusions.

```
data menarche;
input ID Age Age at Men Time r men Perc BF;
datalines;
             1 9.32
                                    -3.87
                                             7.94
                            13.19
             1
                  10.33
                           13.19
                                    -2.86
                                             15.65
             1
                 11.24
                         13.19
                                  -1.95
                                             13.51
                   16.76
                         12.63
                                  4.13 18.53
           162
;
ods rtf file="C:\Users\mclaina\Dropbox\Classes\755 Spring 2018\Examples\14 - Model
Diagnostics.rtf";
ods html;
ods graphics on;
proc mixed data = menarche2;
class ID;
model Perc BF = some model/ solution vciry residual outp=BF pred outpm=BF pred2
influence(effect=ID est);
random some random effects;
run;
```

Model Information							
Data Set	WORK.MENARCHE2						
Dependent Variable	Perc_BF						
<b>Covariance Structure</b>	Unstructured						
Subject Effect	ID						
<b>Estimation Method</b>	REML						
Residual Variance Method	Profile						
Fixed Effects SE Method	Model-Based						
<b>Degrees of Freedom Method</b>	Containment						

Class Level Information						
Class	Levels	Values				
ID	162	1 2 162				

Dimensions	
<b>Covariance Parameters</b>	7
Columns in X	3
Columns in Z per Subject	3
Subjects	162
Max Obs per Subject	10

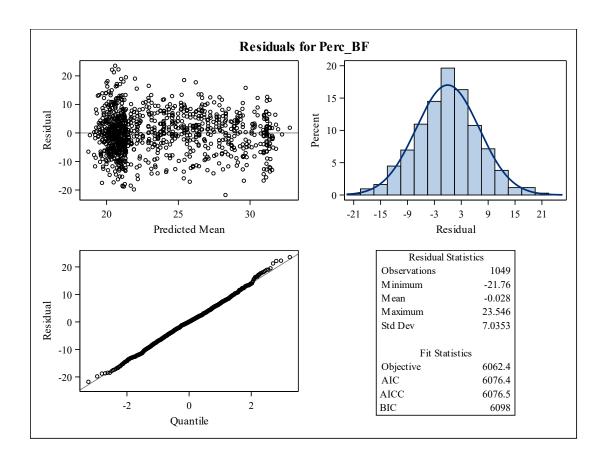
Number of Observations					
<b>Number of Observations Read</b>	1049				
<b>Number of Observations Used</b>	1049				
<b>Number of Observations Not Used</b>	0				

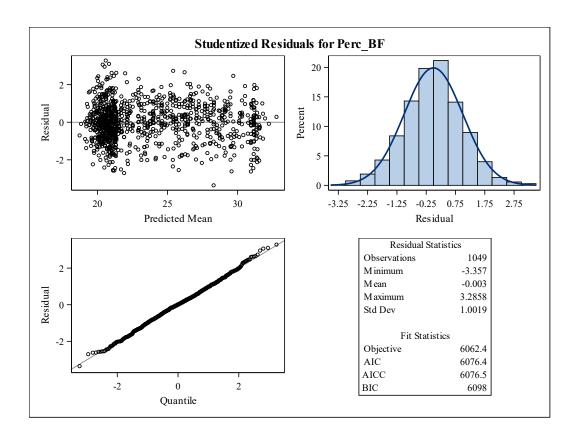
Iteration History									
Iteration	Evaluations	-2 Res Log Like	Criterion						
0	1	7073.41750695							
1	2	6063.49976979	0.00045902						
2	1	6062.44105637	0.00001857						
3	1	6062.40145176	0.00000004						
4	1	6062.40136458	0.00000000						

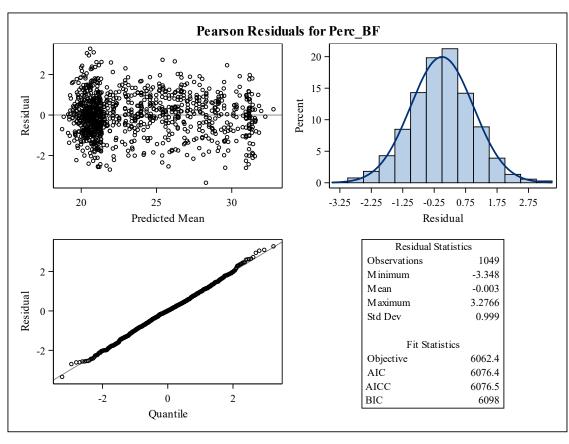
Convergence criteria met.

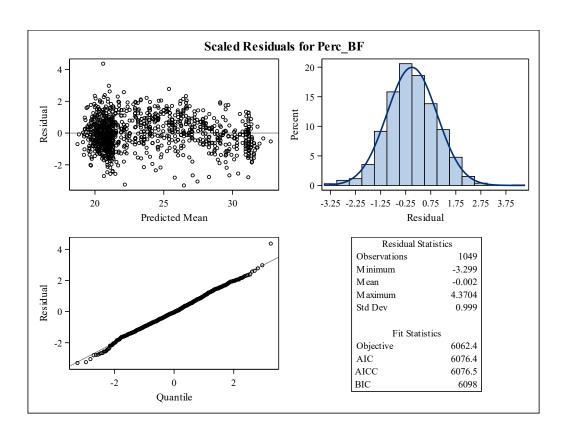
	Estimated V Matrix for ID 1										
Row	Col1	Col2	Col3	Col4	Col5	Col6					
1	60.2897	46.9926	43.5473	39.9506	36.0086	32.8875					
2	46.9926	54.3058	42.8865	40.8548	38.5541	35.3123					
3	43.5473	42.8865	51.7642	41.6694	40.8475	37.4971					
4	39.9506	40.8548	41.6694	51.9930	43.2418	39.7779					
5	36.0086	38.5541	40.8475	43.2418	55.0584	42.0457					
6	32.8875	35.3123	37.4971	39.7779	42.0457	48.8593					

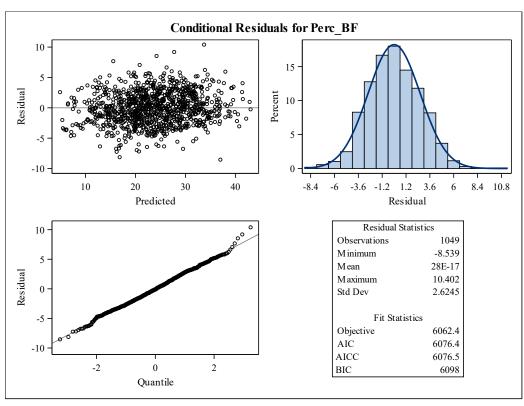
Estimated V Correlation Matrix for ID 1										
Row	Col1	Col2	Col3	Col4	Col5	Col6				
1	1.0000	0.8213	0.7795	0.7136	0.6250	0.6059				
2	0.8213	1.0000	0.8089	0.7689	0.7051	0.6855				
3	0.7795	0.8089	1.0000	0.8032	0.7651	0.7456				
4	0.7136	0.7689	0.8032	1.0000	0.8082	0.7892				
5	0.6250	0.7051	0.7651	0.8082	1.0000	0.8107				
6	0.6059	0.6855	0.7456	0.7892	0.8107	1.0000				

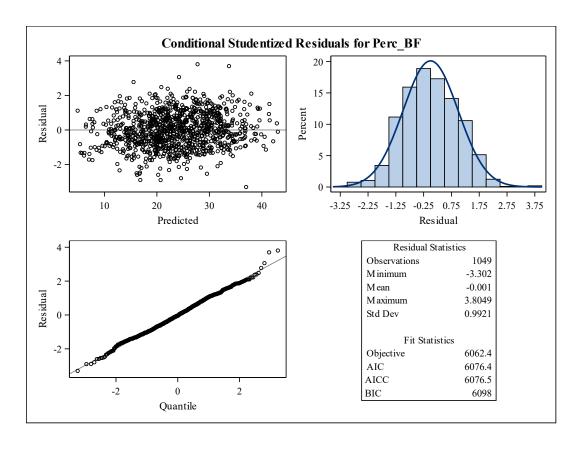


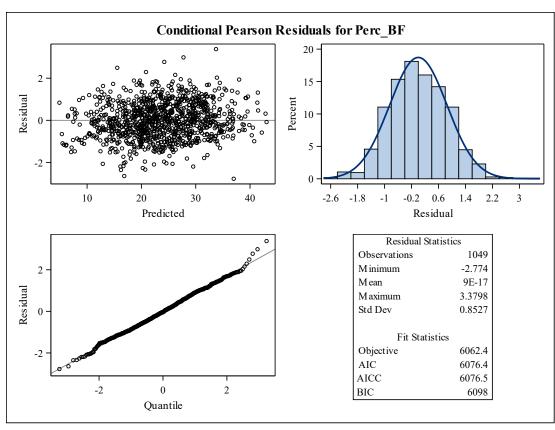




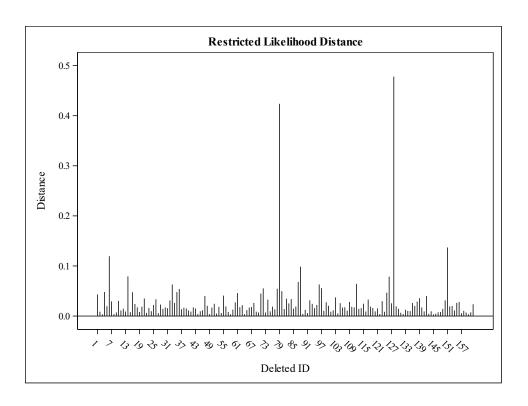


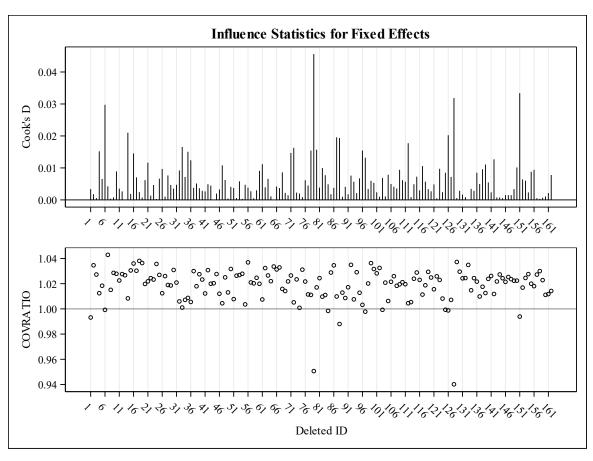


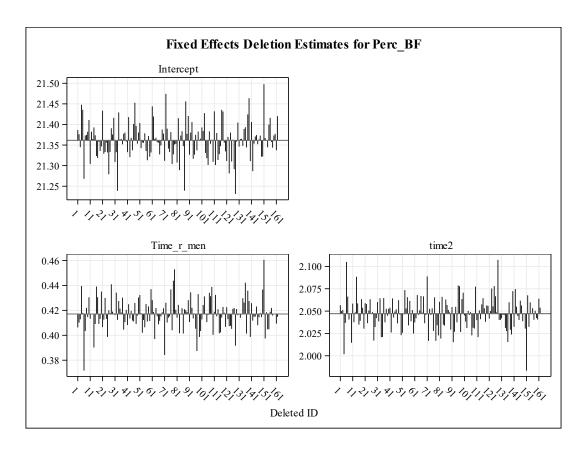


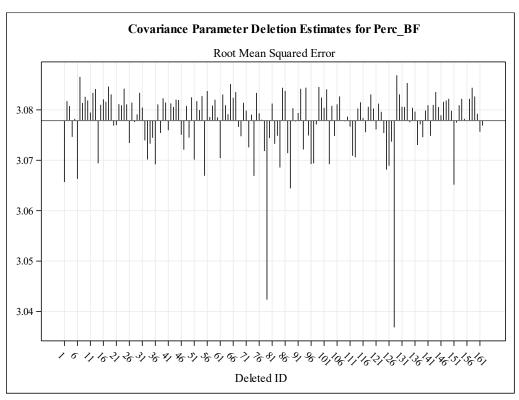


	Influence Diagnostics for Levels of ID										
ID	Number of Observations in Level	PRESS Statistic	Cook's D	MDFFITS	COVRATIO	COVTRACE	RMSE without deleted level	Restricted Likelihood Distance			
1	6	350.35	0.00337	0.00337	0.9932	0.0068	3.06568	0.0432			
2	9	144.73	0.00179	0.00177	1.0346	0.0342	3.08173	0.0087			
3	8	87.53	0.00057	0.00057	1.0273	0.0270	3.08075	0.0036			
4	6	474.26	0.01520	0.01513	1.0124	0.0124	3.07465	0.0480			
5	7	687.09	0.00655	0.00650	1.0184	0.0183	3.07822	0.0197			
161	5	76.23	0.00214	0.00213	1.0118	0.0117	3.07565	0.0075			
162	5	569.33	0.00779	0.00774	1.0142	0.0142	3.07688	0.0236			









```
*Let's take a look at some on the influential diagnostic datasets;

proc print data = BF_pred (obs=5);

run;

proc print data = BF_pred2 (obs=5);

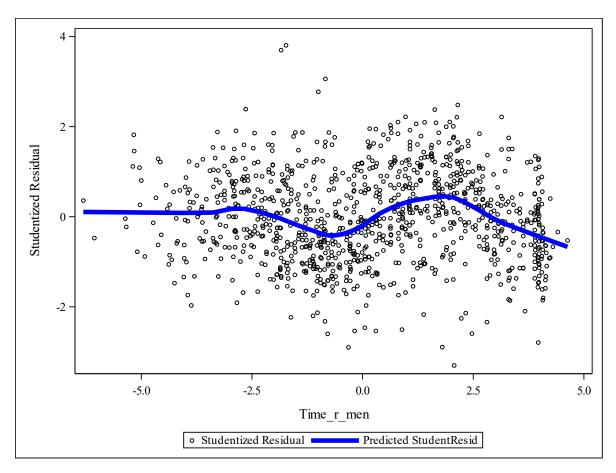
run;
```

								StudentR	PearsonR	
Obs	ID	Pred	StdErrPred	DF	Lower	Upper	Resid	esid	esid	Alpha
1	1	12.4633	2.04879	565	8.4392	16.4875	-4.52333	-1.96933	-1.46964	0.05
2	1	13.5505	1.53271	565	10.5400	16.5610	2.09953	0.78661	0.68214	0.05
3	1	14.5300	1.27459	565	12.0265	17.0335	-1.01998	-0.36408	-0.33139	0.05
4	1	15.5525	1.34887	565	12.9031	18.2020	7.67746	2.77511	2.49442	0.05
5	1	16.7745	1.74260	565	13.3517	20.1973	-6.25451	-2.46528	-2.03210	0.05

											StudentResi	PearsonRes
Obs	ID	Pred	StdErrPred	DF	Alpha	Lower	Upper	Resid	ScaledResid	ScaledDep	d	id
1	1	19.7472	0.66886	565	0.05	18.4334	21.0609	-11.8072	-1.52063	1.02258	-1.52630	-1.52063
2	1	20.1684	0.58520	565	0.05	19.0190	21.3179	-4.5184	1.11420	2.25027	-0.61509	-0.61315
3	1	20.5480	0.53952	565	0.05	19.4883	21.6077	-7.0380	-0.22144	0.75212	-0.98098	-0.97822
4	1	20.9443	0.53017	565	0.05	19.9029	21.9856	2.2857	1.97969	2.79848	0.31786	0.31699
5	1	21.4846	0.56147	565	0.05	20.3818	22.5874	-10.9646	-2.56764	-1.90242	-1.48193	-1.47768

```
*Now let's plot the studentized residuals by time relative to menarche to view any
patterns;
*First, we'll create a loess smooth plot;
proc loess data=BF_pred plots=none;
ods output outputstatistics=out_BF_pred;
model StudentResid=Time_r_men;
run;

proc sort data=out_BF_pred;
by Time_r_men DepVar;
run;
*Now we'll view the loess smooth line by the actual values;
proc sgplot data=out_BF_pred;
scatter x=Time_r_men y=DepVar/ markerattrs=(color=black size=5);
series x=Time_r_men y=pred/ lineattrs=(color=blue thickness=5);
run;
```



\*Lastly let's check the normality of the residuals using the scaled version;

proc univariate data=BF\_pred2 normal;

var ScaledResid;

histogram ScaledResid;

qqplot ScaledResid;

run;

Moments							
N	1049	Sum Weights	1049				
Mean	-0.0022715	<b>Sum Observations</b>	-2.3827516				
<b>Std Deviation</b>	0.99904276	Variance	0.99808644				
Skewness	0.0262867	Kurtosis	0.28659264				
<b>Uncorrected SS</b>	1046	Corrected SS	1045.99459				
Coeff Variation	-43982.589	Std Error Mean	0.03084582				

Basic Statistical Measures						
Loca	tion	Variability				
Mean	-0.00227	<b>Std Deviation</b>	0.99904			
Median	-0.04586	Variance	0.99809			
Mode		Range	7.66893			
		Interquartile Range	1.33801			

Tests for Location: Mu0=0							
Test Statistic p Value							
Student's t	nt's t t		<b>Pr</b> >  t	0.9413			
Sign M		-14.5	Pr >=  M	0.3873			
Signed Rank	S	-2419.5	Pr >=  S	0.8054			

Tests for Normality						
Test	Statistic p Value					
Shapiro-Wilk	W	0.997273	Pr < W	0.0730		
Kolmogorov-Smirnov	D	0.02304	Pr > D	>0.1500		
Cramer-von Mises	W-Sq	0.071804	Pr > W-Sq	>0.2500		
Anderson-Darling	A-Sq	0.501229	Pr > A-Sq	0.2153		

```
*This dataset we'll look at more form influential observations;
data dental;
input Person Gender $ Y1 Y2 Y3 Y4;
y=y1; Age=8; output;
y=y2; Age=10; output;
y=y3; Age=12; output;
y=y4; Age=14; output;
datalines;
                    F
                           21.0
                                    20.0 21.5 23.0
           2
                    F
                            21.0
                                    21.5
                                             24.0
                                                     25.5
           3
                    F
                            20.5
                                    24.0
                                             24.5
                                                     26.0
                                          25.0
                  F
           4
                           23.5 24.5
                                                    26.5
           5
                  F
                           21.5 23.0 22.5
                                                    23.5
           6
                  F
                           20.0
                                   21.0 21.0
                                                    22.5
           7
                  F
                           21.5
                                   22.5 23.0
                                                    25.0
           8
                  F
                           23.0 23.0 23.5
                                                    24.0
                           20.0 21.0 22.0
16.5 19.0 19.0
           9
                   F
                                                    21.5
          10
                   F
                                                    19.5
          11
                  F
                           24.5 25.0 28.0
                                                   28.0
          12
                           26.0 25.0 29.0
                  M
                                                    31.0
          13
                           21.5 22.5 23.0
                  M
                                                  26.5
                           23.0 22.5 24.0
          14
                  M
                                                  27.5
                           25.5 27.5 26.5
20.0 23.5 22.5
24.5 25.5 27.0
22.0 22.0 24.5
24.0 21.5 24.5
          15
                  M
                                                    27.0
                                                    26.0
          16
                  M
          17
                   M
                                                   28.5
                  M
          18
                                                   26.5
          19
                                                  25.5
                  M
          20
                  M
                           23.0 20.5 31.0 26.0

    27.5
    28.0
    31.0

    23.0
    23.0
    23.5

    21.5
    23.5
    24.0

    17.0
    24.5
    26.0

                  M
                                                    31.5
          21
                                                    25.0
          22
                  M
```

run;

23

24

25

26

27

;

M

M

M

M

\*A quick note on the "iter" command. It is the maximum number of additional iterations PROC MIXED performs to update the fixed effects and covariance parameter estimates following data point removal. When n > 0, the influence analysis is iterative. The default is n = 0. If n > 0 and METHOD=REML (default) or METHOD=ML, the procedure updates fixed effects and variance-covariance parameters after removing the selected observations with additional Newton-Raphson iterations, starting from the converged estimates for the entire data. The process stops for each observation or set of observations if the convergence criterion is satisfied or the number of further iterations exceeds n.;

22.5 25.5 25.5

23.0 24.5 26.0

22.0 21.5 23.5 25.0

28.0

29.5

26.0

30.0

```
ods html;
ods graphics on;
proc mixed data=dental;
class Person Gender;
model y = Gender Age Gender*Age / influence(iter=5 effect=Person est);
repeated / type=ar(1) subject=Person;
run;
ods graphics off;
ods html close;
```

Model Information					
Data Set	WORK.DENTAL				
Dependent Variable	У				
Covariance Structure	Autoregressive				
Subject Effect	Person				
Estimation Method	REML				
Residual Variance Method	Profile				
Fixed Effects SE Method	Model-Based				
Degrees of Freedom Method	Between-Within				

Class Level Information						
Class	Levels	Values				
Person	27	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27				
Gender	2	FM				

Dimensions					
Covariance Parameters	2				
Columns in X	6				
Columns in Z	0				
Subjects	27				
Max Obs per Subject	4				

Number of Observations				
Number of Observations Read	108			
Number of Observations Used	108			
Number of Observations Not Used	0			

Covariance Parameter Estimates							
Cov Parm Subject Estimate							
AR(1)	Person	0.6245					
Residual		5.2145					

Fit Statistics					
-2 Res Log Likelihood	444.6				
AIC (Smaller is Better)	448.6				
AICC (Smaller is Better)	448.7				
BIC (Smaller is Better)	451.2				

Null Model Likelihood Ratio Test					
DF Chi-Square Pr > ChiS					
1	38.97	<.0001			

Type 3 Tests of Fixed Effects								
Num Den DF DF F Value Pr > F								
Gender	1	25	0.11	0.7376				
Age	1	79	46.77	<.0001				
Age*Gender	1	79	2.43	0.1233				

	Influence Diagnostics for Levels of Person								
Person	Number of Observations in Level	Iterations	PRESS Statistic	Cook's D	MDFFITS	COVRATIO	COVTRACE	Cook's D CovParms	MDFFITS CovParms
1	4	1	9.6412	0.01119	0.00992	1.3415	0.3098	0.03565	0.0326
2	4	1	3.7181	0.01154	0.01016	1.3815	0.3415	0.05100	0.0456
3	4	1	10.8479	0.02907	0.02591	1.3174	0.2903	0.01812	0.0165
4	4	2	24.4013	0.04667	0.04347	1.1961	0.1888	0.05982	0.0534
5	4	1	1.6900	0.00334	0.00294	1.4051	0.3600	0.06207	0.0548
6	4	1	11.7185	0.01981	0.01776	1.3158	0.2891	0.05000	0.0463
7	4	1	1.2187	0.00307	0.00269	1.4069	0.3613	0.06890	0.0611
8	4	1	5.0250	0.01807	0.01592	1.3675	0.3305	0.05455	0.0495
9	4	1	13.1216	0.03196	0.02875	1.2864	0.2649	0.03419	0.0319
10	4	2	85.2778	0.16899	0.18880	0.7277	0.2899	0.61657	0.6538
11	4	2	69.6436	0.12270	0.12869	0.8583	0.1389	0.32291	0.3103
12	4	2	38.8782	0.05476	0.05435	0.9723	0.0252	0.04632	0.0488
13	4	1	14.8373	0.01148	0.01060	1.2248	0.2102	0.02479	0.0231
14	4	1	6.8024	0.00082	0.00074	1.2851	0.2613	0.03085	0.0272
15	4	1	20.5090	0.03478	0.03244	1.1510	0.1454	0.00823	0.0077
16	4	1	23.5720	0.02813	0.02629	1.1361	0.1319	0.00310	0.0030
17	4	1	8.1930	0.01197	0.01103	1.2538	0.2350	0.05914	0.0545
18	4	1	9.1204	0.00677	0.00619	1.2680	0.2468	0.04130	0.0377
19	4	2	15.6824	0.02364	0.02219	1.1953	0.1854	0.03370	0.0347
20	4	4	42.9162	0.00831	0.00825	0.8175	0.0354	7.42313	19.6236
21	4	2	91.5142	0.12647	0.14980	0.6399	0.4057	0.82320	0.8390
22	4	1	14.2947	0.02470	0.02276	1.2119	0.1991	0.02903	0.0271
23	4	1	6.4046	0.00940	0.00859	1.2817	0.2584	0.02976	0.0263
24	4	2	42.5667	0.14725	0.15055	0.8947	0.1059	0.48613	0.5319
25	4	1	3.6903	0.00450	0.00409	1.2990	0.2727	0.03991	0.0351
26	4	1	7.9353	0.01924	0.01766	1.2430	0.2258	0.01989	0.0181
27	4	2	21.6103	0.02450	0.02298	1.1631	0.1564	0.02558	0.0237

Influence Diagnostics for Levels of Person								
Person	COVRATIO CovParms	COVTRACE CovParms	RMSE without deleted level	Restricted Likelihood Distance				
1	1.1489	0.1448	2.31316	0.078				
2	1.1493	0.1469	2.32185	0.093				
3	1.1232	0.1211	2.30818	0.132				
4	1.1237	0.1234	2.27905	0.248				
5	1.1579	0.1560	2.32689	0.072				
6	1.1510	0.1458	2.30704	0.127				
7	1.1609	0.1583	2.32712	0.077				
8	1.1499	0.1464	2.31862	0.122				
9	1.1269	0.1232	2.30052	0.160				
10	0.9747	0.0896	2.14317	1.492				
11	1.0449	0.1108	2.18660	0.899				
12	1.0277	0.0361	2.23808	0.275				
13	1.1387	0.1345	2.30444	0.070				
14	1.1352	0.1351	2.31932	0.032				
15	1.1126	0.1098	2.28639	0.145				
16	1.0687	0.0677	2.28347	0.114				
17	1.1559	0.1505	2.31089	0.103				
18	1.1495	0.1455	2.31460	0.066				
19	1.0665	0.0702	2.29951	0.129				
20	0.5307	0.1129	2.27626	14.546				
21	1.0053	0.1676	2.12644	1.463				
22	1.1312	0.1273	2.30118	0.126				
23	1.1284	0.1285	2.31849	0.064				
24	0.8505	0.1552	2.21905	1.170				
25	1.1378	0.1376	2.32227	0.055				
26	1.1154	0.1140	2.30915	0.095				
27	1.1264	0.1232	2.28908	0.125				

