

SAS/IML[®] Software: Changes and Enhancements, Release 8.1

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SAS/IML® Software: Changes and Enhancements, Release 8.1

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Chapter 1

Changes and Enhancements

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Chapter 1

Changes and Enhancements

Overview

New financial functions, robust regression subroutines, and subroutines related to multivariate time series analysis have been added. All necessary details such as arguments and operands are included.

Financial Functions

The following functions that compute information about financial quantities have been added:

CONVEXIT Function

calculates and returns a scalar containing the convexity of a noncontingent cash flow

CONVEXIT(*times,flows,ytm*)

The CONVEXIT function calculates and returns a scalar containing the convexity of a noncontingent cash flow.

<i>times</i>	is an n -dimensional column vector of times. Elements should be non-negative.
<i>flows</i>	is an n -dimensional column vector of cash flows.
<i>ytm</i>	is the per-period yield-to-maturity of the cash-flow stream. This is a scalar and should be positive.

Convexity is essentially a measure of how duration, the sensitivity of price to yield, changes as interest rates change:

$$C = \frac{1}{P} \frac{d^2 P}{dy^2}$$

With cash flows that are not yield-sensitive, and the assumption of parallel shifts to a flat term-structure, convexity is given by

$$C = \frac{\sum_{k=1}^K t_k (t_k + 1) \frac{c(k)}{(1+y)^{t_k}}}{P(1+y)^2}$$

where P is the present value, y is the effective per period yield-to-maturity, K is the number of cash flows, and the k -th cash flow is $c(k)$ t_k periods from the present.

The statements

```
timesn=T(do(1,100,1));
flows=repeat(10,100);
ytm=0.1;
convexit=convexit(timesn,flows,ytm);
print convexit;
```

result in the following output:

```
CONVEXIT
199.26229
```

DURATION Function

calculates and returns a scalar containing the modified duration of a noncontingent cash flow

DURATION(*times,flows,ytm*)

The DURATION function returns the modified duration of a noncontingent cash flow as a scalar.

<i>times</i>	is an n -dimensional column vector of times. Elements should be non-negative.
<i>flows</i>	is an n -dimensional column vector of cash flows.
<i>ytm</i>	is the per-period yield-to-maturity of the cash-flow stream. This is a scalar and should be positive.

Duration of a security is generally defined as

$$D = -\frac{\frac{dP}{P}}{dy}$$

In other words, it is the relative change in price for a unit change in yield. Since prices move in the opposite direction to yields, the sign change preserves positivity for convenience. With cash flows that are not yield-sensitive, and the assumption of parallel shifts to a flat term-structure, duration is given by

$$D_{\text{mod}} = \frac{\sum_{k=1}^K t_k \frac{c(k)}{(1+y)^{t_k}}}{P(1+y)}$$

where P is the present value, y is the per period effective yield-to-maturity, K is the number of cash flows, and the k -th cash flow is $c(k)$, t_k periods from the present. This

measure is referred to as *modified duration* to differentiate it from the first duration measure ever proposed, *Macauley duration*:

$$D_{\text{Mac}} = \frac{\sum_{k=1}^K t_k \frac{c(k)}{(1+y)^{t_k}}}{P}$$

This expression also reveals the reason for the name duration, since it is a present-value-weighted average of the duration (that is, timing) of all the cash flows and is hence an “average time-to-maturity” of the bond.

For example, the statements below

```
times={1};
ytm={0.1};
flow={10};
duration=duration(times,flow,ytm);
print duration;
```

produce the output

```
DURATION
0.9090909
```

FORWARD Function

calculates a column vector of forward rates given vectors of spot rates and times

FORWARD(*times*,*spot_rates*)

The FORWARD function returns an $n \times 1$ vector of forward rates.

times is an $n \times 1$ column vector of times in consistent units. Elements should be non-negative.

spot_rates is an $n \times 1$ column vector of corresponding per-period spot rates. Elements should be positive.

The FORWARD function transforms the given spot rates as

$$f_1 = s_1$$

$$f_i = \left(\frac{(1 + s_i)^{t_i}}{(1 + s_{i-1})^{t_{i-1}}} \right)^{\frac{1}{t_i - t_{i-1}}} - 1; \quad i = 2, \dots, n$$

For example, the following statements

```

spt={0.75};
times={1};
forward=forward(times,spt);
print forward;

```

produce the following output:

```

FORWARD
0.75

```

PV Function

calculates the present value of a vector of cash flows and returns a scalar

PV(times,flows,freq,rates)

The PV function returns a scalar containing the present value of the cash flows based on the specified frequency and rates.

times	is an $n \times 1$ column vector of times. Elements should be non-negative.
flows	is an $n \times 1$ column vector of cash flows.
freq	is a scalar that represents the base of the rates to be used for discounting the cash flows. If positive, it represents discrete compounding as the reciprocal of the number of compoundings. If zero, it represents continuous compounding. If -1, it represents per-period discount factors. No other negative values are allowed.
rates	is an $n \times 1$ column vector of rates to be used for discounting the cash flows. Elements should be positive.

A general present value relationship can be written as

$$P = \sum_{k=1}^K c(k)D(t_k)$$

where P is the present value of the asset, $\{c(k)\}_{k=1, \dots, K}$ is the sequence of cash flows from the asset, t_k is the time to the k -th cash flow in periods from the present, and $D(t)$ is the discount function for time t .

With per-unit-time-period discount factors d_t :

$$D(t) = d_t^t$$

With continuous compounding:

$$D(t) = e^{-r_t t}$$

With discrete compounding:

$$D(t) = (1 + fr)^{-t/f}$$

where $f > 0$ is the frequency, the reciprocal of the number of compoundings per unit time period.

The following code presents an example of the PV function:

```
timesn=T(do(1,100,1));
flows=repeat(10,100);
freq=50;
rate=repeat(0.10,100);
pv=pv(timesn,flows,freq,rate);
print pv;
```

The result is

```
PV
266.4717
```

RATES Function

calculates a column vector of interest rates converted from one base to another

RATES(rates,oldfreq,newfreq)

The RATES function returns an $n \times 1$ vector of interest rates converted from one base to another.

rates	is an $n \times 1$ column vector of rates. Elements should be positive.
oldfreq	is a scalar that represents the old base. If positive, it represents discrete compounding as the reciprocal of the number of compoundings. If zero, it represents continuous compounding. If -1, it represents discount factors. No other negative values are allowed.
newfreq	is a scalar that represents the new base. If positive, it represents discrete compounding as the reciprocal of the number of compoundings. If zero, it represents continuous compounding. If -1, it represents per-period discount factors. No other negative values are allowed.

Let $D(t)$ be the discount function, which is the present value of a unit amount to be received t periods from now. The discount function can be expressed in three different ways:

with per-unit-time-period discount factors d_t :

$$D(t) = d_t^t$$

with continuous compounding:

$$D(t) = e^{-r_t t}$$

with discrete compounding:

$$D(t) = (1 + fr)^{-t/f}$$

where $f > 0$ is the frequency, the reciprocal of the number of compoundings per unit time period. The RATES function converts between these three representations.

For example, the following code uses the RATES function:

```
rates=T(do(0.1,0.3,0.1));
oldfreq=0;
newfreq=0;
rates=rates(rates,oldfreq,newfreq);
print rates;
```

The output is

```
RATES
0.1
0.2
0.3
```

SPOT Function

calculates a column vector of spot rates given vectors of forward rates and times

SPOT(times,forward_rates)

The SPOT function returns an $n \times 1$ vector of spot rates.

times is an $n \times 1$ column vector of times in consistent units. Elements should be non-negative.

forward_rates is an $n \times 1$ column vector of corresponding per-period forward rates. Elements should be positive.

The SPOT function transforms the given spot rates as

$$s_1 = f_1$$

$$s_i = \left(\prod_{j=1}^{j=i} (1 + f_j)^{t_j - t_{j-1}} \right)^{\frac{1}{t_i}} - 1; \quad i = 2, \dots, n$$

where, by convention, $t_0 = 0$.

For example, the following code

```
fwd={0.05};
times={1};
spot=spot(times,fwd);
print spot;
```

produces the following output:

```
SPOT
0.05
```

YIELD Function

calculates yield-to-maturity of a cash-flow stream and returns a scalar

YIELD(*times,flows,freq,value*)

The YIELD function returns a scalar containing yield-to-maturity of a cash flow stream based on frequency and value specified.

<i>times</i>	is an n -dimensional column vector of times. Elements should be non-negative.
<i>flows</i>	is an n -dimensional column vector of cash flows.
<i>freq</i>	is a scalar that represents the base of the rates to be used for discounting the cash flows. If positive, it represents discrete compounding as the reciprocal of the number of compoundings. If zero, it represents continuous compounding. No negative values are allowed.
<i>value</i>	is a scalar that is the discounted present value of the cash flows.

The present value relationship can be written as

$$P = \sum_{k=1}^K c(k)D(t_k)$$

where P is the present value of the asset, $\{c(k)\}_{k=1, \dots, K}$ is the sequence of cash flows from the asset, t_k is the time to the k -th cash flow in periods from the present, and $D(t)$ is the discount function for time t .

With continuous compounding:

$$D(t) = e^{-yt}$$

With discrete compounding:

$$D(t) = (1 + fy)^{-t/f}$$

where $f > 0$ is the frequency, the reciprocal of the number of compoundings per unit time period, and y is the yield-to-maturity. The YIELD function solves for y .

For example, the following code

```
timesn=T(do(1,100,1));
flows=repeat(10,100);
freq=50;
value=682.31027;
yield=yield(timesn,flows,freq,value);
print yield;
```

produces the following output:

```
YIELD
0.0100001
```

Robust Regression

These are the new algorithms for robust regression analysis:

LTS Call

performs robust regression

```
CALL LTS(sc, coef, wgt, opt, y <, < x ><, sorb>>);
```

A new algorithm, FAST-LTS, was added to the LTS subroutine in SAS/IML Release 8.1. The FAST-LTS algorithm is set as the default algorithm. The original algorithm is kept for convenience, and can be used temporarily by specifying optn[9]=1. Eventually the original algorithm will be replaced with the new FAST-LTS algorithm.

The original algorithm for the LTS subroutine and the algorithm used in the LMS subroutine are based on the PROGRESS program by Rousseeuw and Leroy (1987). Rousseeuw and Hubert (1996) prepared a new version of PROGRESS to facilitate its inclusion in SAS software, and they have incorporated several recent developments. Among other things, the new version of PROGRESS now yields the exact LMS for simple regression, and the program uses a new definition of the robust coefficient of determination (R^2). Therefore, the outputs may differ slightly from those given in Rousseeuw and Leroy (1987) or those obtained from software based on the older version of PROGRESS.

FAST-LTS

Least trimmed squares (LTS) regression is based on the subset of h cases (out of n) whose least squares fit possesses the smallest sum of squared residuals. The coverage h may be set between $n/2$ and n . The LTS method was proposed by Rousseeuw (1984, p. 876) as a highly robust regression estimator, with breakdown value $(n-h)/n$. It turned out that the computation time of the previous LTS algorithm grew too fast with the size of the data set, precluding their use for data mining. Rousseeuw and

Van Driessen (1998) developed a new algorithm called FAST-LTS. The basic idea is an inequality involving order statistics and sums of squared residuals. Based on this inequality, techniques called “selective iteration” and “nested extensions” are developed in Rousseeuw and Van Driessen (1998). The new LTS algorithm implements these techniques to achieve faster computation. The intercept adjustment technique is also used in this new algorithm. For small data sets, FAST-LTS typically finds the exact LTS, whereas for larger data sets it gives more accurate results than the previous LTS algorithm and is faster by orders of magnitude. The new algorithm is described briefly as follows; refer to Rousseeuw and Van Driessen (1998) for details:

1. The default h is $(n + p + 1)/2$, where p is the number of the independent variables. You can choose any integer h with $[(n + p + 1)/2] \leq h \leq n$. The LTS’s breakdown point $(n - h + 1)/n$ is reported. If you are sure that the data contains less than 25% of contamination, you can obtain a good compromise between breakdown value and statistical efficiency by putting $h = \lceil .75n \rceil$.
2. If $p = 1$ (univariate data), then compute the LTS estimator by the exact algorithm of Rousseeuw and Leroy (1987, pp. 171-172) and stop.
3. From here on, $p \geq 2$. If $n < 600$, draw a random p -subset and compute the regression coefficients using these p points (if the regression is degenerate, draw another p -subset). Compute the absolute residuals for all points in the data set and select the first h points with smallest absolute residuals. From this selected h -subset, carry out C-steps (Concentration step; refer to Rousseeuw and Van Driessen [1998] for details) until convergence. Repeat this procedure 500 or $\binom{n}{p}$ times (as determined by $opt[5]$ of LTS Call) and find the ten (at most) solutions with the lowest sums of h squared residuals. For each of these ten best solutions, take C-steps until convergence and find the best final solution.
4. If $n > 600$, construct up to five disjoint random subsets with sizes as equal as possible, but not to exceed 300. Inside each subset, repeat the procedure in step 3 $500/5 = 100$ times and keep the ten best solutions. Pool the subsets, yielding the merged set of size n_{merged} . In the merged set, for each of the $5 \times 10 = 50$ best solutions, carry out two C-steps using n_{merged} and $h_{\text{merged}} = \lceil n_{\text{merged}}(h/n) \rceil$ and keep the ten best solutions. In the full data set, for each of these ten best solutions, take C-steps using n and h until convergence and find the best final solution.

OPTION Changes

The previous LTS algorithm is used if $optn[9] = 1$; the FAST-LTS algorithm is set as default (or $optn[9] = 0$).

OUTPUT Changes

Because of the change in algorithm, the output from the FAST-LTS algorithm is different:

1. The “Complete Enumeration for LTS” table and the “Resistant Diagnostic” table do not apply for the FAST-LTS algorithm and are not displayed.
2. The analysis based on “Observations of Best Subset” of size p , where p is the number of the independent variables, is replaced by the analysis based on

observations of the best h of the entire data set obtained after full iteration, which gives the exact LTS estimator and covariances for small data set ($\binom{n}{p} < 500$). LTS Objective Function, Preliminary LTS Scale, Robust R Squared, and Final LTS Scale are also reported for the LTS estimator.

3. The LTS residuals are changed, because of the change of the LTS estimator.
4. The “Coef” vector does not include the best subset.

See the following illustrative example for details.

Illustrative Example

The following example shows the difference between the two algorithms. The second output is generated by the FAST-LTS algorithm.

```

title1 'Compare two algorithms for LTS';

proc iml;
  reset noname;

  x = {42, 37, 37, 28, 18, 18, 19, 20, 15};
  y = {80, 80, 75, 62, 62, 62, 62, 62, 58};
  optn = j(9, 1, .);
  optn[1]= 0;      /* --- with intercept      --- */
  optn[2]= 4;      /* --- print all output      --- */
  optn[3]= 3;      /* --- compute LS and WLS    --- */
  optn[8]= 3;      /* --- covariance matrices   --- */
  optn[9]= 1;      /* --- Version 7 LTS         --- */

  call lts(sc, coef, wgt, optn, y, x);

  print "sc is " sc, "coef is " coef;

  optn[9]= 0;      /* --- FAST-LTS algorithm    --- */

  call lts(sc, coef, wgt, optn, y, x);

  print "sc is " sc, "coef is " coef;

quit;

```

Comparison of the outputs is summarized as follows:

1. The summary statistics for dependent and independent variables and the results of the classical (unweighted) least-squares estimation (outputs in Page 1 and first half of Page 2) do not change.
2. The “Complete Enumeration for LTS” table on page 1 and the “Resistant Diagnostic” table on page 3 of the first output is eliminated. The analysis based on “Observations of Best Subset” of size p , where p is the number of the independent variables, is replaced by the analysis based on observations of the best h of the entire data set obtained after full iteration.

3. The best ten (at most) estimates before the final search are available by setting a proper option (see pages 2 and 3 of the second output). The first one is the best with the smallest objective value.
4. The results of the weighted least squared estimation (page 4 for the first output, page 5 for the second output) do not change because the two algorithms detect the same outliers in this simple example. Results will be changed if they detect different outliers.
5. The “Coef” vector does not include the best subset (page 6 of the second output).

Compare two algorithms for LTS

2

LS Residuals

N	Observed	Estimated	Residual	Res / S
1	80.000000	79.880383	0.119617	0.038891
2	80.000000	75.855263	4.144737	1.347581
3	75.000000	75.855263	-0.855263	-0.278072
4	62.000000	68.610048	-6.610048	-2.149130
5	62.000000	60.559809	1.440191	0.468251
6	62.000000	60.559809	1.440191	0.468251
7	62.000000	61.364833	0.635167	0.206512
8	62.000000	62.169856	-0.169856	-0.055226
9	58.000000	58.144737	-0.144737	-0.047058

Distribution of Residuals

MinRes	1st Qu.	Median
-6.610047847	-0.512559809	0.1196172249
Mean	3rd Qu.	MaxRes
-2.36848E-15	1.0376794258	4.1447368421

(no change)

There are 36 subsets of 2 cases out of 9 cases.

All 36 subsets will be considered.

Complete Enumeration for LTS

Subset	Singular	Best Criterion	Percent
10	1	0.084493	27
18	1	0.084493	50
28	2	0.084493	77
36	2	0.084493	100

Minimum Criterion= 0.0844927545

(eliminated)

Least Trimmed Squares (LTS) Method
 Minimizing Sum of 6 Smallest Squared Residuals.
 Highest Possible Breakdown Value = 44.44 %
 Selection of All 36 Subsets of 2 Cases Out of 9
 Among 36 subsets 2 are singular.

Observations of Best Subset

1 5

Estimated Coefficients

VAR1	Intercep
0.75	47.916666667

Compare two algorithms for LTS

3

LTS Objective Function = 0.6236095645

Preliminary LTS Scale = 1.189465671

Robust R Squared = 0.825

Final LTS Scale = 0.8595864639

LTS Residuals

N	Observed	Estimated	Residual	Res / S
1	80.000000	79.416667	0.583333	0.678621
2	80.000000	75.666667	4.333333	5.041184
3	75.000000	75.666667	-0.666667	-0.775567
4	62.000000	68.916667	-6.916667	-8.046505
5	62.000000	61.416667	0.583333	0.678621
6	62.000000	61.416667	0.583333	0.678621
7	62.000000	62.166667	-0.166667	-0.193892
8	62.000000	62.916667	-0.916667	-1.066404
9	58.000000	59.166667	-1.166667	-1.357242

Distribution of Residuals

MinRes	1st Qu.	Median
-6.916666667	-1.041666667	-0.166666667
Mean	3rd Qu.	MaxRes
-0.416666667	0.583333333	4.333333333

(replaced)

Resistant Diagnostic

N	U	Resistant Diagnostic
1	12.521981	5.600000
2	12.521981	5.600000
3	9.167879	4.100000
4	13.339459	5.965588
5	1.709204	0.764379
6	1.709204	0.764379
7	0.642081	0.287147
8	1.697749	0.759257
9	2.236068	1.000000

Median(U)= 2.2360679775

(eliminated)

Compare two algorithms for LTS

4

Weighted Least-Squares Estimation

RLS Parameter Estimates Based on LTS

Variable	Estimate	Approx Std Err	t Value	Pr > t	Lower WCI	Upper WCI
VAR1	0.76455907	0.03125286	24.46	<.0001	0.70330458	0.82581356
Intercep	47.3985025	0.815574	58.12	<.0001	45.8000068	48.9969982

Weighted Sum of Squares = 3.3544093178

Degrees of Freedom = 5

RLS Scale Estimate = 0.819073784

Cov Matrix of Parameter Estimates

	VAR1	Intercep
VAR1	0.0009767415	-0.02358133
Intercep	-0.02358133	0.6651609492

Weighted R-squared = 0.9917145853

F(1,5) Statistic = 598.47009637

Probability = 2.1279132E-6

There are 7 points with nonzero weight.

Average Weight = 0.7777777778

Weighted LS Residuals

N	Observed	Estimated	Residual	Res / S	Weight
1	80.000000	79.509983	0.490017	0.598257	1.000000
2	80.000000	75.687188	4.312812	5.265474	0
3	75.000000	75.687188	-0.687188	-0.838982	1.000000
4	62.000000	68.806156	-6.806156	-8.309577	0
5	62.000000	61.160566	0.839434	1.024858	1.000000
6	62.000000	61.160566	0.839434	1.024858	1.000000
7	62.000000	61.925125	0.074875	0.091414	1.000000
8	62.000000	62.689684	-0.689684	-0.842029	1.000000
9	58.000000	58.866889	-0.866889	-1.058377	1.000000

Distribution of Residuals

MinRes	1st Qu.	Median
-6.806156406	-0.77828619	0.074875208
Mean	3rd Qu.	MaxRes
-0.27703827	0.6647254576	4.31281198

The run has been executed successfully.

(no change)

Compare two algorithms for LTS

5

```

sc is      6
           36
           2
           7
           0.6236096
           1.1894657
           0.8595865
           0.825
           1.9073884
           .
           0.8190738

```

```

           3.3544093
           0.9917146
           598.4701
           .
           .
           .
           .
           .
           .

```

```

coef is      0.75 47.916667
              1      5 ----->eliminated
0.7645591 47.398502
0.0312529 0.815574
24.463648 58.11674
2.1279E-6 2.8538E-8
0.7033046 45.800007
0.8258136 48.996998

```


Compare two algorithms for LTS

2

LS Residuals

N	Observed	Estimated	Residual	Res / S
1	80.000000	79.880383	0.119617	0.038891
2	80.000000	75.855263	4.144737	1.347581
3	75.000000	75.855263	-0.855263	-0.278072
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7	62.000000	61.364833	0.635167	0.206512
8	62.000000	62.169856	-0.169856	-0.055226
9	58.000000	58.144737	-0.144737	-0.047058

Distribution of Residuals

MinRes	1st Qu.	Median
-6.610047847	-0.512559809	0.1196172249
Mean	3rd Qu.	MaxRes
-2.36848E-15	1.0376794258	4.1447368421

(no change)

Least Trimmed Squares (LTS) Method

The (at most) 10 Best Estimates

Objective Value [1]: 0.0428203092

Estimated Coefficients

VAR1	Intercep
0.7521545304	0.1250675364

Objective Value [2]: 0.0454534796

Estimated Coefficients

VAR1	Intercep
0.7773132092	0.0679381212

Objective Value [3]: 0.0458503276

Compare two algorithms for LTS

3

Estimated Coefficients

VAR1	Intercep
0.7857569344	0.0875053435

Objective Value [4]: 0.0470161862

Estimated Coefficients

VAR1	Intercep
0.7454450208	0.1033087106

Objective Value [5]: 0.0504563846

Estimated Coefficients

VAR1	Intercep
0.987904817	0.1606655488

Objective Value [6]: 0.1790484378

Estimated Coefficients

VAR1	Intercep
0.1926222834	-0.081665104

Objective Value [7]: 1.797693E308

(added)

Least Trimmed Squares (LTS) Method
 Minimizing Sum of 6 Smallest Squared Residuals.
 Highest Possible Breakdown Value = 44.44 %
 Selection of All 36 Subsets of 2 Cases Out of 9
 Among 36 subsets 2 are singular.

```

Compare two algorithms for LTS
4

The best half of the entire data set obtained after full iteration consists of
the cases:

      1      3      5      6      7      8

Estimated Coefficients

      VAR1      Intercep
      0.7488687783      47.945701357

LTS Objective Function = 0.6235087791

Preliminary LTS Scale = 1.1892734341

Robust R Squared = 0.819730444

Final LTS Scale = 0.8627851118

LTS Residuals

N      Observed      Estimated      Residual      Res / S
1      80.000000      79.398190      0.601810      0.697520
2      80.000000      75.653846      4.346154      5.037354
3      75.000000      75.653846      -0.653846      -0.757832
4      62.000000      68.914027      -6.914027      -8.013614
5      62.000000      61.425339      0.574661      0.666053
6      62.000000      61.425339      0.574661      0.666053
7      62.000000      62.174208      -0.174208      -0.201914
8      62.000000      62.923077      -0.923077      -1.069880
9      58.000000      59.178733      -1.178733      -1.366195

Distribution of Residuals

      MinRes      1st Qu.      Median
      -6.914027149      -1.050904977      -0.174208145

      Mean      3rd Qu.      MaxRes
      -0.416289593      0.5746606335      4.3461538462

(changed)
-----

```

Compare two algorithms for LTS

5

Weighted Least-Squares Estimation

RLS Parameter Estimates Based on LTS

Variable	Estimate	Approx Std Err	t Value	Pr > t	Lower WCI	Upper WCI
VAR1	0.76455907	0.03125286	24.46	<.0001	0.70330458	0.82581356
Intercep	47.3985025	0.815574	58.12	<.0001	45.8000068	48.9969982

Weighted Sum of Squares = 3.3544093178

Degrees of Freedom = 5

RLS Scale Estimate = 0.819073784

Cov Matrix of Parameter Estimates

	VAR1	Intercep
VAR1	0.0009767415	-0.02358133
Intercep	-0.02358133	0.6651609492

Weighted R-squared = 0.9917145853

F(1,5) Statistic = 598.47009637

Probability = 2.1279132E-6

There are 7 points with nonzero weight.

Average Weight = 0.7777777778

Weighted LS Residuals

N	Observed	Estimated	Residual	Res / S	Weight
1	80.000000	79.509983	0.490017	0.598257	1.000000
2	80.000000	75.687188	4.312812	5.265474	0
3	75.000000	75.687188	-0.687188	-0.838982	1.000000
4	62.000000	68.806156	-6.806156	-8.309577	0
5	62.000000	61.160566	0.839434	1.024858	1.000000
6	62.000000	61.160566	0.839434	1.024858	1.000000
7	62.000000	61.925125	0.074875	0.091414	1.000000
8	62.000000	62.689684	-0.689684	-0.842029	1.000000
9	58.000000	58.866889	-0.866889	-1.058377	1.000000

Distribution of Residuals

MinRes	1st Qu.	Median
-6.806156406	-0.77828619	0.074875208
Mean	3rd Qu.	MaxRes
-0.27703827	0.6647254576	4.31281198

The run has been executed successfully.

(no change)

```

Compare two algorithms for LTS
6

sc is      6
           36
           2
           7
           0.6235088
           1.1892734
           0.8627851
           0.8197304
           1.879
           .
           0.8190738
           3.3544093
           0.9917146
           598.4701
           .
           .
           .
           .
           .

coef is 0.7488688 47.945701
        0.7645591 47.398502
        0.0312529 0.815574
        24.463648 58.11674
        2.1279E-6 2.8538E-8
        0.7033046 45.800007
        0.8258136 48.996998
        .         .

                                         (changed)
-----

```

MCD Call

finds the minimum covariance determinant estimator

CALL MCD(*sc*, *coef*, *dist*, *opt*, *x*);

The MCD call is the robust (resistant) estimation of multivariate location and scatter, defined by minimizing the determinant of the covariance matrix computed from h points. The algorithm for the MCD subroutine is based on the FAST-MCD algorithm given by Rousseeuw and Van Driessen (1999).

The MCD subroutine computes the minimum covariance determinant estimator. These robust locations and covariance matrices can be used to detect multivariate outliers and leverage points. For this purpose, the MCD subroutine provides a table of robust distances.

In the following discussion, N is the number of observations and n is the number of regressors. The inputs to the MCD subroutine are as follows:

opt refers to an options vector with the following components (missing values

are treated as default values):

opt[1] specifies the amount of printed output. Higher option values request additional output and include the output of lower values.

opt[1]=0 prints no output except error messages.

opt[1]=1 prints most of the output.

opt[1]=2 additionally prints case numbers of the observations in the best subset and some basic history of the optimization process.

opt[1]=3 additionally prints how many subsets result in singular linear systems.

The default is *opt*[1]=0.

opt[2] specifies whether the classical, initial, and final robust covariance matrices are printed. The default is *opt*[2]=0. Note that the final robust covariance matrix is always returned in *coef*.

opt[3] specifies whether the classical, initial, and final robust correlation matrices are printed or returned:

opt[3]=0 does not return or print.

opt[3]=1 prints the robust correlation matrix.

opt[3]=2 returns the final robust correlation matrix in *coef*.

opt[3]=3 prints and returns the final robust correlation matrix.

opt[4] specifies the quantile h used in the objective function. The default is *opt*[5]= $h = \lceil \frac{N+n+1}{2} \rceil$. If the value of h is specified outside the range $\frac{N}{2} + 1 \leq h \leq \frac{3N}{4} + \frac{n+1}{4}$, it is reset to the closest boundary of this region.

opt[5] specifies the number N_{Rep} of subset generations. This option is the same as described for the LTS subroutines. Due to computer time restrictions, not all subset combinations can be inspected for larger values of N and n .

When *opt*[5] is zero or missing:

If $N > 600$, construct up to five disjoint random subsets with sizes as equal as possible, but not to exceed 300. Inside each subset, choose $500/5 = 100$ subset combinations of n observations.

If $N < 600$, the number of subsets is taken from the following table.

n	1	2	3	4	5	6	7	8	9	10
N_{lower}	500	50	22	17	15	14	0	0	0	0

n	11	12	13	14	15
N_{lower}	0	0	0	0	0

If the number of cases (observations) N is smaller than N_{lower} , then all possible subsets are used; otherwise, 500 subsets are cho-

sen randomly. This means that an exhaustive search is performed for $opt[5]=-1$. If N is larger than N_{upper} , a note is printed in the log file indicating how many subsets exist.

x refers to an $N \times n$ matrix \mathbf{X} of regressors.

Missing values are not permitted in x . Missing values in opt cause the default value to be used.

The MCD subroutine returns the following values:

sc is a column vector containing the following scalar information:

$sc[1]$	the quantile h used in the objective function
$sc[2]$	number of subsets generated
$sc[3]$	number of subsets with singular linear systems
$sc[4]$	number of nonzero weights w_i
$sc[5]$	lowest value of the objective function F_{MCD} attained (smallest determinant)
$sc[6]$	Mahalanobis-like distance used in the computation of the lowest value of the objective function F_{MCD}
$sc[7]$	the cutoff value used for the outlier decision

$coef$ is a matrix with n columns containing the following results in its rows:

$coef[1]$	location of ellipsoid center
$coef[2]$	eigenvalues of final robust scatter matrix
$coef[3:2+n]$	the final robust scatter matrix for $opt[2]=1$ or $opt[2]=3$
$coef[2+n+1:2+2n]$	the final robust correlation matrix for $opt[3]=1$ or $opt[3]=3$

$dist$ is a matrix with N columns containing the following results in its rows:

$dist[1]$	Mahalanobis distances
$dist[2]$	robust distances based on the final estimates
$dist[3]$	weights (=1 for small, =0 for large robust distances)

Example

Consider Brownlee's (1965) stackloss data used in the example for the MVE subroutine.

For $N = 21$ and $n = 4$ (three explanatory variables including intercept), you obtain a total of 5,985 different subsets of 4 observations out of 21. If you decide not to specify $optn[5]$, the MCD algorithm chooses 500 random sample subsets:

```

/* X1  X2  X3   Y      Stackloss data */
aa = { 1  80  27  89  42,
        1  80  27  88  37,
        1  75  25  90  37,
        1  62  24  87  28,
        1  62  22  87  18,
        1  62  23  87  18,
        1  62  24  93  19,
        1  62  24  93  20,
        1  58  23  87  15,
        1  58  18  80  14,
        1  58  18  89  14,
        1  58  17  88  13,
        1  58  18  82  11,
        1  58  19  93  12,
        1  50  18  89   8,
        1  50  18  86   7,
        1  50  19  72   8,
        1  50  19  79   8,
        1  50  20  80   9,
        1  56  20  82  15,
        1  70  20  91  15 };

a = aa[,2:4];
optn = j(8,1,.);
optn[1]= 2;          /* ipri */
optn[2]= 1;          /* pcov: print COV */
optn[3]= 1;          /* pcov: print CORR */

CALL MCD(sc,xmcd,dist,optn,a);

```

The first part of the output of this program is a summary of the MCD algorithm and the final h points selected:

```

Fast MCD by Rousseeuw and Van Driessen

Number of Variables                3
Number of Observations             21
Default Value for h                12
Specified Value for h              12
Breakdown Value                    42.86
- Highest Possible Breakdown Value -

```

The best half of the entire data set obtained after full iteration consists of the cases:

```

4      5      6      7      8      9      10     11     12     13     14     20

```

The second part of the output is the MCD estimators of the location, scatter matrix, and correlation matrix:

MCD Location Estimate

VAR1	VAR2	VAR3
59.5	20.833333333	87.333333333
Average of 12 Selected Points		

MCD Scatter Matrix Estimate

	VAR1	VAR2	VAR3
VAR1	5.1818181818	4.8181818182	4.7272727273
VAR2	4.8181818182	7.6060606061	5.0606060606
VAR3	4.7272727273	5.0606060606	19.151515152

Determinant = 238.07387929

Covariance Matrix of 12 Selected Points

MCD Correlation Matrix

	VAR1	VAR2	VAR3
VAR1	1	0.7674714142	0.4745347313
VAR2	0.7674714142	1	0.4192963398
VAR3	0.4745347313	0.4192963398	1

The MCD scatter matrix is multiplied by a factor to make it consistent when all the data come from a single Gaussian distribution.

Consistent Scatter Matrix

	VAR1	VAR2	VAR3
VAR1	8.6578437815	8.0502757968	7.8983838007
VAR2	8.0502757968	12.708297013	8.4553211199
VAR3	7.8983838007	8.4553211199	31.998580526
Determinant = 397.77668436			

The final output presents a table containing the classical Mahalanobis distances, the robust distances, and the weights identifying the outlying observations (that is, leverage points when explaining y with these three regressor variables):

Classical Distances and Robust (Rousseeuw) Distances			
Unsquared Mahalanobis Distance and			
Unsquared Rousseeuw Distance of Each Observation			
N	Mahalanobis Distances	Robust Distances	Weight
1	2.253603	12.173282	0
2	2.324745	12.255677	0
3	1.593712	9.263990	0
4	1.271898	1.401368	1.000000

5	0.303357	1.420020	1.000000
6	0.772895	1.291188	1.000000
7	1.852661	1.460370	1.000000
8	1.852661	1.460370	1.000000
9	1.360622	2.120590	1.000000
10	1.745997	1.809708	1.000000
11	1.465702	1.362278	1.000000
12	1.841504	1.667437	1.000000
13	1.482649	1.416724	1.000000
14	1.778785	1.988240	1.000000
15	1.690241	5.874858	0
16	1.291934	5.606157	0
17	2.700016	6.133319	0
18	1.503155	5.760432	0
19	1.593221	6.156248	0
20	0.807054	2.172300	1.000000
21	2.176761	7.622769	0

Robust distances are based on reweighted estimates.

The cutoff value is the square root of the 0.975 quantile of the chi square distribution with 3 degrees of freedom.

Points whose robust distance exceeds 3.0575159206 have received a zero weight in the last column above.

There were 9 such points in the data.

These may include boundary cases.

Only points whose robust distance is substantially larger than the cutoff should be considered outliers.

Multivariate Time Series Analysis

These are the new functions for the creation and analysis of multivariate time series:

VARMACOV Call

computes the theoretical cross-covariance matrices for a stationary VARMA(p, q) model

CALL VARMACOV(*cov, phi, theta, sigma* <, *p, q, lag*>);

The inputs to the VARMACOV subroutine are as follows:

phi specifies a $km_p \times k$ matrix, Φ , containing the autoregressive coefficient matrices, where m_p is the number of elements in the subset of the AR order and $k \geq 2$ is the number of variables. All the roots of $|\Phi(B)| = 0$ should be greater than one in absolute value, where $\Phi(B)$ is the finite order matrix polynomial in the backshift operator B , such that $B^j \mathbf{y}_t = \mathbf{y}_{t-j}$. You must

specify either *phi* or *theta*.

- theta* specifies a $km_q \times k$ matrix containing the moving-average coefficient matrices, where m_q is the number of the elements in the subset of the MA order. You must specify either *phi* or *theta*.
- sigma* specifies a $k \times k$ symmetric positive-definite covariance matrix of the innovation series. If *sigma* is not specified, then an identity matrix is used.
- p* specifies the subset of the AR order. The quantity m_p is defined as

$$m_p = \text{nrow}(\text{phi}) / \text{ncol}(\text{phi})$$

where $\text{nrow}(\text{phi})$ is the number of rows of the matrix *phi* and $\text{ncol}(\text{phi})$ is the number of columns of the matrix *phi*.

If you do not specify *p*, the default subset is $p = \{1, 2, \dots, m_p\}$.

For example, consider a four-dimensional vector time series, and *phi* is a 4×4 matrix. If you specify $p=1$ (the default, since $m_p = 4/4 = 1$), the VARMA(1) subroutine computes the theoretical cross-covariance matrices of VAR(1) as $\mathbf{y}_t = \Phi \mathbf{y}_{t-1} + \epsilon_t$.

If you specify $p=2$, the VARMA(2) subroutine computes the cross-covariance matrices of VAR(2) as $\mathbf{y}_t = \Phi \mathbf{y}_{t-2} + \epsilon_t$.

Let $\text{phi} = [\Phi'_1, \Phi'_2]'$ be an 8×4 matrix. If you specify $p = \{1, 3\}$, the VARMA(3) subroutine computes the cross-covariance matrices of VAR(3) as $\mathbf{y}_t = \Phi_1 \mathbf{y}_{t-1} + \Phi_2 \mathbf{y}_{t-3} + \epsilon_t$. If you do not specify *p*, the VARMA(2) subroutine computes the cross-covariance matrices of VAR(2) as $\mathbf{y}_t = \Phi_1 \mathbf{y}_{t-1} + \Phi_2 \mathbf{y}_{t-2} + \epsilon_t$.

- q* specifies the subset of the MA order. The quantity m_q is defined as

$$m_q = \text{nrow}(\text{theta}) / \text{ncol}(\text{theta})$$

where $\text{nrow}(\text{theta})$ is the number of rows of matrix *theta* and $\text{ncol}(\text{theta})$ is the number of columns of matrix *theta*.

If you do not specify *q*, the default subset is $q = \{1, 2, \dots, m_q\}$.

The usage of *q* is the same as that of *p*.

- lag* specifies the length of lags, which must be a positive number. If $\text{lag} = h$, the VARMA(1,1) computes the cross-covariance matrices from lag zero to lag *h*. By default, $\text{lag} = 12$.

The VARMA(1,1) subroutine returns the following value:

- cov* is a $k(\text{lag} + 1) \times k$ matrix that contains the theoretical cross-covariance matrices of the VARMA(*p*, *q*) model.

To compute the cross-covariance matrices of a bivariate ($k = 2$) VARMA(1,1) model

$$\mathbf{y}_t = \Phi \mathbf{y}_{t-1} + \epsilon_t - \Theta \epsilon_{t-1}$$

where

$$\Phi = \begin{bmatrix} 1.2 & -0.5 \\ 0.6 & 0.3 \end{bmatrix} \quad \Theta = \begin{bmatrix} -0.6 & 0.3 \\ 0.3 & 0.6 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1.0 & 0.5 \\ 0.5 & 1.25 \end{bmatrix}$$

you can specify

```
phi = { 1.2 -0.5, 0.6 0.3 };
theta= {-0.6 0.3, 0.3 0.6 };
sigma= { 1.0 0.5, 0.5 1.25};
call varmacov(cov, phi, theta, sigma) lag=5;
```

VARMALIK Call

computes the log-likelihood function for a VARMA(p, q) model

CALL VARMALIK(*lnl, series, phi, theta, sigma* <, *p, q, opt*>);

The inputs to the VARMALIK subroutine are as follows:

- | | |
|---------------|--|
| <i>series</i> | specifies an $n \times k$ matrix containing the vector time series (assuming mean zero), where n is the number of observations and $k \geq 2$ is the number of variables. |
| <i>phi</i> | specifies a $km_p \times k$ matrix containing the autoregressive coefficient matrices, where m_p is the number of the elements in the subset of the AR order. You must specify either <i>phi</i> or <i>theta</i> . |
| <i>theta</i> | specifies a $km_q \times k$ matrix containing the moving-average coefficient matrices, where m_q is the number of the elements in the subset of the MA order. You must specify either <i>phi</i> or <i>theta</i> . |
| <i>sigma</i> | specifies a $k \times k$ covariance matrix of the innovation series. If you do not specify <i>sigma</i> , an identity matrix is used. |
| <i>p</i> | specifies the subset of the AR order. See the VARMACOV subroutine. |
| <i>q</i> | specifies the subset of the MA order. See the VARMACOV subroutine. |
| <i>opt</i> | specifies the method of computing the log-likelihood function: |
| <i>opt</i> =0 | requests the multivariate innovations algorithm. This algorithm requires that the time series is stationary and does not contain missing observations. |
| <i>opt</i> =1 | requests the conditional log-likelihood function. This algorithm requires that the number of the observations in the time series must be greater than $p+q$ and that the series does not contain missing observations. |
| <i>opt</i> =2 | requests the Kalman filtering algorithm. This is the default and is used if the required conditions in <i>opt</i> =0 and <i>opt</i> =1 are not satisfied. |

The VARMA1IK subroutine returns the following value:

lnl is a 3×1 matrix containing the log-likelihood function, the sum of log determinant of the innovation variance, and the weighted sum of squares of residuals. The log-likelihood function is computed as $-0.5 \times$ (the sum of last two terms).

The options $opt=0$ and $opt=2$ are equivalent for stationary time series without missing values. Setting $opt=0$ is useful for a small number of the observations and a high order of p and q ; $opt=1$ is useful for a high order of p and q ; $opt=2$ is useful for a low order of p and q , or for missing values in the observations.

To compute the log-likelihood function of a bivariate ($k = 2$) VARMA(1,1) model

$$\mathbf{y}_t = \Phi \mathbf{y}_{t-1} + \epsilon_t - \Theta \epsilon_{t-1}$$

where

$$\Phi = \begin{bmatrix} 1.2 & -0.5 \\ 0.6 & 0.3 \end{bmatrix} \quad \Theta = \begin{bmatrix} -0.6 & 0.3 \\ 0.3 & 0.6 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1.0 & 0.5 \\ 0.5 & 1.25 \end{bmatrix}$$

you can specify

```
phi = { 1.2 -0.5, 0.6 0.3 };
theta= {-0.6 0.3, 0.3 0.6 };
sigma= { 1.0 0.5, 0.5 1.25};
call varmasim(yt, phi, theta) sigma=sigma;
call varmalik(lnl, yt, phi, theta, sigma);
```

VARMASIM Call

generates a VARMA(p,q) time series

CALL VARMA SIM(*series, phi, theta, mu, sigma, n <, p, q, initial, seed>*);

The inputs to the VARMA SIM subroutine are as follows:

- phi* specifies a $km_p \times k$ matrix containing the autoregressive coefficient matrices, where m_p is the number of the elements in the subset of the AR order and $k \geq 2$ is the number of variables. You must specify either *phi* or *theta*.
- theta* specifies a $km_q \times k$ matrix containing the moving-average coefficient matrices, where m_q is the number of the elements in the subset of the MA order. You must specify either *phi* or *theta*.
- mu* specifies a $k \times 1$ (or $1 \times k$) mean vector of the series. If *mu* is not specified, a zero vector is used.
- sigma* specifies a $k \times k$ covariance matrix of the innovation series. If *sigma* is not specified, an identity matrix is used.

- n* specifies the length of the series. If *n* is not specified, $n = 100$ is used.
- p* specifies the subset of the AR order. See the VARMA COV subroutine.
- q* specifies the subset of the MA order. See the VARMA COV subroutine.
- initial* specifies the initial values of random variables. If *initial* = a_0 , then $\mathbf{y}_{-p+1}, \dots, \mathbf{y}_0$ and $\boldsymbol{\epsilon}_{-q+1}, \dots, \boldsymbol{\epsilon}_0$ all take the same value a_0 . If the *initial* option is not specified, the initial values are estimated for the stationary vector time series; the initial values are assumed as zero for the nonstationary vector time series.
- seed* specifies the random number seed. See the VNORMAL subroutine.

The VARMA SIM subroutine returns the following value:

- series* is an $n \times k$ matrix containing the generated VARMA(p, q) time series. When either the *initial* option is specified or zero initial values are used, these initial values are not included in *series*.

To generate a bivariate ($k = 2$) stationary VARMA(1,1) time series

$$\mathbf{y}_t - \boldsymbol{\mu} = \Phi(\mathbf{y}_{t-1} - \boldsymbol{\mu}) + \boldsymbol{\epsilon}_t - \Theta\boldsymbol{\epsilon}_{t-1}$$

where

$$\Phi = \begin{bmatrix} 1.2 & -0.5 \\ 0.6 & 0.3 \end{bmatrix} \quad \Theta = \begin{bmatrix} -0.6 & 0.3 \\ 0.3 & 0.6 \end{bmatrix} \quad \boldsymbol{\mu} = \begin{bmatrix} 10 \\ 20 \end{bmatrix} \quad \Sigma = \begin{bmatrix} 1.0 & 0.5 \\ 0.5 & 1.25 \end{bmatrix}$$

you can specify

```
phi = { 1.2 -0.5, 0.6 0.3 };
theta= {-0.6 0.3, 0.3 0.6 };
mu = { 10, 20 };
sigma= { 1.0 0.5, 0.5 1.25};
call varmasim(yt, phi, theta, mu, sigma, 100);
```

To generate a bivariate ($k = 2$) nonstationary VARMA(1,1) time series with the same $\boldsymbol{\mu}$, Σ , and Θ in the previous example and the AR coefficient

$$\Phi = \begin{bmatrix} 1.0 & 0 \\ 0 & 0.3 \end{bmatrix}$$

you can specify

```
phi = { 1.0 0.0, 0.0 0.3 };
call varmasim(yt, phi, theta, mu, sigma, 100) initial=3;
```

VNORMAL Call

generates a multivariate normal random series

CALL VNORMAL(*series*, *mu*, *sigma*, *n* <, *seed*>);

The inputs to the VNORMAL subroutine are as follows:

- mu* specifies a $k \times 1$ (or $1 \times k$) mean vector, where $k \geq 2$ is the number of variables. You must specify either *mu* or *sigma*. If *mu* is not specified, a zero vector is used.
- sigma* specifies a $k \times k$ symmetric positive-definite covariance matrix. By default, *sigma* is an identity matrix with dimension k . You must specify either *mu* or *sigma*. If *sigma* is not specified, an identity matrix is used.
- n* specifies the length of the series. If *n* is not specified, $n = 100$ is used.
- seed* specifies the random number seed. If it is not supplied, the system clock is used to generate the seed. If it is negative, then the absolute value is used as the starting seed; otherwise, subsequent calls ignore the value of *seed* and use the last seed generated internally.

The VNORMAL subroutine returns the following value:

series is an $n \times k$ matrix that contains the generated normal random series.

To generate a bivariate ($k = 2$) normal random series with mean μ and covariance matrix Σ , where

$$\mu = \begin{bmatrix} 10 \\ 20 \end{bmatrix} \quad \text{and} \quad \Sigma = \begin{bmatrix} 1.0 & 0.5 \\ 0.5 & 1.25 \end{bmatrix}$$

you can specify

```
mu      = { 10, 20 };
sigma= { 1.0  0.5, 0.5 1.25};
call vnormal(et, mu, sigma, 100);
```

VTSROOT Call

calculates the characteristic roots of the model from AR and MA characteristic functions

CALL VTSROOT(*root*, *phi*, *theta*<, *p*, *q*>);

The inputs to the VTSROOT subroutine are as follows:

- phi* specifies a $km_p \times k$ matrix containing the autoregressive coefficient matrices, where m_p is the number of the elements in the subset of the AR order and $k \geq 2$ is the number of variables. You must specify either *phi* or *theta*.
- theta* specifies a $km_q \times k$ matrix containing the moving-average coefficient matrices, where m_q is the number of the elements in the subset of the MA order. You must specify either *phi* or *theta*.
- p* specifies the subset of the AR order. See the VARMA COV subroutine.
- q* specifies the subset of the MA order. See the VARMA COV subroutine.

The VTSROOT subroutine returns the following value:

root is a $k(p_{max} + q_{max}) \times 5$ matrix, where p_{max} is the maximum order of the AR characteristic function and q_{max} is the maximum order of the MA characteristic function. The first kp_{max} rows refer to the results of the AR characteristic function; the last q_{max} rows refer to the results of the MA characteristic function.

The first column contains the real parts, x , of eigenvalues of companion matrix associated with the AR(p_{max}) or MA(q_{max}) characteristic function; the second column contains the imaginary parts, y , of the eigenvalues; the third column contains the moduli of the eigenvalues, $\sqrt{x^2 + y^2}$; the fourth column contains the arguments ($\arctan(y/x)$) of the eigenvalues, measured in radians from the positive real axis. The fifth column contains the arguments expressed in degrees rather than radians.

To compute the roots of the characteristic functions, $\Phi(B) = I - \Phi B$ and $\Theta(B) = I - \Theta B$, where I is an identity matrix with dimension 2 and

$$\Phi = \begin{bmatrix} 1.2 & -0.5 \\ 0.6 & 0.3 \end{bmatrix} \quad \Theta = \begin{bmatrix} -0.6 & 0.3 \\ 0.3 & 0.6 \end{bmatrix}$$

you can specify

```
phi = { 1.2 -0.5, 0.6 0.3 };
theta= {-0.6 0.3, 0.3 0.6 };
call vtsroot(root, phi, theta);
```

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