

Simulating with Parameter Uncertainty

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1 Purpose

This script shows how to conduct a simulation that considers uncertainty in the parameter estimates.

2 Data

Here we load metrumrg and read in the data to be used for simulations.

Listing 1:

```
> library(metrumrg)
> data <- read.csv("../data/derived/phase1.csv")</pre>
> head(data)
 C ID TIME SEQ EVID AMT
                          DV SUBJ HOUR TAFD TAD LDOS MDV HEIGHT WEIGHT SEX
1 C 1 0.00 0
                 0
                                1 0.00 0.00
                                                           174
                                                                 74.2
2 . 1 0.00 1
                 1 1000
                                1 0.00 0.00
                                              0 1000
                                                                 74.2
3 . 1 0.25 0
                 0
                      . 0.363
                                1 0.25 0.25 0.25 1000 0
                                                           174
                                                                74.2
                                                                       0
4 . 1 0.50 0
                      . 0.914
                                1 0.50 0.50 0.5 1000
                                                                74.2
                                                           174
                                                                       0
                      . 1.12
5 . 1 1.00 0
                                1 1.00 1.00
                                              1 1000 0
                                                                74.2
                                                           174
6 . 1 2.00
                      . 2.28
                                              2 1000 0
                                                           174 74.2 0
                 0
                                1 2.00 2.00
  AGE DOSE FED SMK DS CRCN predose zerodv
1 29.1 1000 1
                0 0 83.5
2 29.1 1000
                0 0 83.5
                               0
                                      0
                0 0 83.5
3 29.1 1000
4 29.1 1000
                0 0 83.5
                               0
                                      0
5 29.1 1000
                0 0 83.5
                               0
                                      0
6 29.1 1000 1 0 0 83.5
                               0
                                      0
```

We use NONMEM output from a simple two compartment model to generate parameters. We use 1005.lst and 1005.cov output from NM7 to populate a call to metrumrg::simpar().



Listing 2:

```
> cov <- read.table("../nonmem/1005/1005.cov", skip=1, header=T)</pre>
> head(cov)
   NAME
             THETA1
                         THETA2
                                      THETA3
                                                  THETA4
                                                              THETA5
1 THETA1 0.8759870 0.79288000 1.06070e-03 0.06301820 -1.7590100
2 THETA2 0.7928800 4.74648000
                               6.68044e-03 0.89631100
                                                           5.2850000
3 THETA3 0.0010607 0.00668044
                                2.75847e-05
                                             0.00222162
                                                          -0.0305675
4 THETA4 0.0630182 0.89631100 2.22162e-03 0.28693100
                                                           0.1902840
5 THETA5 -1.7590100 5.28500000 -3.05675e-02 0.19028400 566.1060000
6 THETA6 -0.0421234 -0.02569100 -1.05405e-04 -0.01064250
                                                           0.7806910
                                                      SIGMA.2.2. OMEGA.1.1.
        THETA6
                             SIGMA.1.1. SIGMA.2.1.
                     THETA7
1 -0.042123400 -0.180383000 -5.30926e-04
                                                  0 2.09474e-02 6.20306e-03
2 -0.025691000 0.066978500 -3.13015e-03
                                                  0 1.92041e-02 5.76993e-03
3 -0.000105405 -0.000138306 -1.02726e-05
                                                  0 5.91125e-05 3.21603e-06
4 -0.010642500 0.015518300 -6.29448e-04
                                                  0 2.54028e-03 4.29694e-03
5 0.780691000 -0.602816000 4.56339e-02
                                                  0 -4.26533e-01 2.73809e-01
                                                  0 -1.07146e-03 1.64777e-03
6 0.013065500 0.000798277 1.21261e-04
   OMEGA.2.1.
                OMEGA.2.2.
                             OMEGA.3.1.
                                          OMEGA.3.2.
                                                        OMEGA.3.3.
1 -1.83821e-04 -4.40561e-03 -5.44206e-03 -2.59575e-03 -3.35118e-03
2 -2.19650e-02 -2.44862e-02 -1.96039e-02 -1.12215e-02 4.77949e-03
3 - 6.50573e - 05 - 7.81727e - 05 - 6.76632e - 05 - 2.75979e - 05 2.83104e - 05
4 -6.21366e-03 -7.78974e-03 -4.55505e-03 -2.25175e-03 3.07354e-03
5 1.60570e-01 2.81746e-02 -4.85574e-03 7.48252e-02 -3.43426e-02
6 3.02269e-04 6.01990e-04 -5.29743e-04 -5.31172e-05 -3.35780e-04
```

We are interested in theta covariance, so we remove extra columns and rows.

Listing 3:

```
> cov <- cov[1:7,c(2:8)]
```



3 Parameters

Now we generate 10 sets of population parameters based on the 1005.lst results.

Listing 4:

```
> set.seed(10)
> PKparms <- simpar(
      nsim=10,
      theta=c(8.58,21.6, 0.0684, 3.78, 107, 0.999, 1.67),
     covar=cov,
     omega=list(0.196, 0.129, 0.107),
     odf=c(40,40,40),
     sigma=list(0.0671),
      sdf=c(200)
> PKparms
    TH.1 TH.2
                  TH.3 TH.4
                              TH.5
                                      TH.6 TH.7 OM1.1
                                                          OM2.2
                                                                  OM3.3
9.472 24.04 0.06312 3.508 106.50 1.0140 1.589 0.1847 0.15400 0.13630
2 10.740 22.97 0.06794 3.814 111.40 0.8452 1.296 0.2862 0.12000 0.16400
3 9.001 21.22 0.06626 3.966 139.60 1.0780 1.512 0.1647 0.12770 0.11300
4 10.680 23.59 0.07180 3.906 121.20 1.1480 1.009 0.1886 0.11460 0.08460
5 10.030 23.02 0.06999 3.573 99.98 0.9437 1.762 0.1526 0.08448 0.13140
6 8.964 21.73 0.06877 3.343 97.72 1.0600 1.806 0.2462 0.17640 0.08805
7 8.630 19.70 0.06611 3.377 135.80 0.9626 1.348 0.2221 0.14440 0.09957
8 9.213 21.23 0.06096 3.082 115.70 1.0420 1.752 0.2287 0.13820 0.06118
9 8.974 23.95 0.06980 4.110 145.70 0.9455 1.782 0.1765 0.12310 0.08504
10 8.939 22.72 0.06436 3.704 113.10 1.0610 1.581 0.2116 0.11940 0.09954
    SG1.1
1 0.06894
2 0.06099
3 0.06041
4 0.07700
```



```
5 0.06269
6 0.07274
7 0.06160
8 0.06692
9 0.06092
10 0.06269
```

4 Control Streams

We read in a control stream and clean out extra xml markup.

Listing 5:

```
> ctl <- as.nmctl(readLines("../nonmem/ctl/1005.ctl"))
> ctl[] <- lapply(ctl,function(rec)sub("<.*","",rec))</pre>
```

Now we iterate across the rows of PKparms, writing out a separate ctl for each.

Listing 6:



5 Simulation

Finally, we run NONMEM simulations using NONR.

Listing 7:

```
> NONR72(
+ run=1:10,
+ command="/opt/NONMEM/nm72/nmqual/autolog.pl",
+ project="../nonmem/sim",
+ diag=FALSE,
+ checkrunno=FALSE,
+ grid=TRUE
+ )
```