

Modeling

April 3, 2012

Tim Bergsma



1 Purpose

This script runs NONMEM models and diagnostics for sample phase1 data.

2 Model Development

2.1 Set up for NONMEM run.

Listing 1:

```
> library(metrumrg)
> #command <- '/opt/NONMEM/nm72/nmqual/autolog.pl'
> cat.cov='SEX'
> cont.cov=c('HEIGHT','WEIGHT','AGE')
> par.list=c('CL','Q','KA','V','V2','V3')
> eta.list=paste('ETA',1:10,sep='')
```

2.2 Run NONMEM.

Listing 2:

```
> NONR72(
+ run=1001:1005,
+ #command=command,
+ project='../nonmem',
+ grid=FALSE,
+ nice=TRUE,
+ checkrunno=FALSE,
+ cont.cov=cont.cov,
+ cat.cov=cat.cov,
+ par.list=par.list,
```



```
+ eta.list=eta.list,
+ plotfile='../nonmem/*/*.pdf',
+ streams='../nonmem/ctl'
+ )
```

Covariance succeeded on model 1005. We can make a quick run log using some simple tools. Table 1.

Listing 3:

```
> log <- rlog(1001:1005,'../nonmem',tool='nm7',file=NULL)
> head(log)
```

	tool	run	parameter	moment	value
1	nm7	1001	ofv	minimum	2526.39867230031
2	nm7	1001	THETA1	estimate	11.7167
3	nm7	1001	THETA1	prse	8.67
4	nm7	1001	THETA1	se	1.01636
5	nm7	1001	THETA2	estimate	14.5657
6	nm7	1001	THETA2	prse	8.67

Listing 4:

> tail(log)

```
tool run parameter
                        moment
245 nm7 1005 SIGMA2.2
                          prse
246 nm7 1005 SIGMA2.2
                            se
247 nm7 1005
                   COV
                        status
248 nm7 1005
                  prob
                          text
249 nm7 1005
                  min
                       status
250 nm7 1005
                 data filename
                                               value
245
                                                33.5
246
                                            0.0676412
247
                                                   0
```



```
248 1005 phase1 2 CMT like 1004 but diff. initial on V3
249
250
                         ../../data/derived/phase1.csv
                                                    Listing 5:
> sapply(log,class)
      tool
                   run parameter
                                       moment
                                                    value
"character" "character" "character" "character"
                                                    Listing 6:
> log$tool <- NULL
> unique(log$parameter)
[1] "ofv"
               "THETA1"
                          "THETA2" "THETA3" "OMEGA1.1" "OMEGA2.1"
[7] "OMEGA2.2" "OMEGA3.1" "OMEGA3.2" "OMEGA3.3" "SIGMA1.1" "SIGMA2.1"
[13] "SIGMA2.2" "cov"
                          "prob"
                                     "min"
                                               "data"
                                                          "THETA4"
[19] "THETA5" "OMEGA4.1" "OMEGA4.2" "OMEGA4.3" "OMEGA4.4" "OMEGA5.1"
[25] "OMEGA5.2" "OMEGA5.3" "OMEGA5.4" "OMEGA5.5" "THETA6" "THETA7"
                                                    Listing 7:
> log <- log[log$parameter %in% c('ofv','prob','cov','min'),]</pre>
> log
    run parameter moment
1 1001
              ofv minimum
38 1001
              cov status
39 1001
           prob
                   text
40 1001
            min status
42 1002
            ofv minimum
112 1002
            cov status
113 1002
            prob
                   text
```



```
114 1002
              min status
116 1003
              ofv minimum
153 1003
            cov status
154 1003
             prob
                   text
155 1003
            min status
157 1004
            ofv minimum
194 1004
            cov status
195 1004
            prob
                     text
196 1004
            min status
198 1005
            ofv minimum
247 1005
            cov status
248 1005
             prob
                    text
249 1005
              min status
                                                        value
1
                                              2526.39867230031
38
39
                                              1001 phase1 1CMT
40
42
                                              2525.96526753388
112
113
                                             1002 phase1 2 CMT
114
116
                                              2569.89393760215
153
154 1003 phase1 2 CMT like 1002 but no eta on Q/v3 and no + err
155
157
                                              2570.45022637547
194
195
                 1004 phase1 2 CMT like 1003 but better bounds
196
198
                                              2405.91625845151
247
248
           1005 phase1 2 CMT like 1004 but diff. initial on V3
249
```



Listing 8:

> with(log, constant(moment,within=parameter))#i.e., moment is non-informative here.

[1] TRUE

Listing 9:

```
> log <- data.frame(cast(log,run~parameter))
> log <- shuffle(log,'prob','run')
> log$ofv <- signif(as.numeric(as.character(log$ofv,6)))</pre>
```

Table 1: Run Log

run	prob	cov	min	ofv
1001	1001 phase1 1CMT	0	0	2526.40
1002	1002 phase1 2 CMT	1	1	2525.97
1003	1003 phase1 2 CMT like 1002 but no eta on Q/v3 and no + err	1	0	2569.89
1004	1004 phase1 2 CMT like 1003 but better bounds	0	0	2570.45
1005	1005 phase1 2 CMT like 1004 but diff. initial on V3	0	0	2405.92

3 Predictive Check

3.1 Create a simulation control stream.

Convert control stream to R object.

Listing 10:

```
> ctl <- read.nmctl('../nonmem/ctl/1005.ctl')</pre>
```



Strip comments and view.

Listing 11:

```
> ctl[] <- lapply(ctl, function(rec) sub(' *; .*', '', rec))</pre>
> ctl
 [1] "$PROB 1005 phase1 2 CMT like 1004 but diff. initial on V3"
 [2] "$INPUT C ID TIME SEQ=DROP EVID AMT DV SUBJ HOUR TAFD TAD LDOS MDV HEIGHT WT SEX AGE DOSE FED"
 [3] "$DATA ../../data/derived/phase1.csv IGNORE=C"
 [4] "$SUBROUTINE ADVAN4 TRANS4"
 [5] "$PK"
 [6] " CL=THETA(1) *EXP(ETA(1)) * THETA(6) **SEX * (WT/70) **THETA(7)"
 [7] " V2 =THETA(2) *EXP(ETA(2))"
 [8] " KA=THETA(3) *EXP(ETA(3))"
[9] " Q =THETA(4)"
[10] " V3=THETA(5)"
[11] " S2=V2"
[12] " "
[13] "$ERROR"
[14] " Y=F*(1+ERR(1)) + ERR(2)"
[15] " IPRE=F"
[16] ""
[17] "$THETA"
[18] "(0,10,50)"
[19] "(0,10,100)"
[20] "(0,0.2, 5)"
[21] "(0,10,50)"
[22] "(0,100,1000)"
[23] "(0,1,2)"
[24] "(0,0.75,3)"
[25] ""
[26] "$OMEGA BLOCK(3)"
[27] ".1"
[28] ".01 .1"
```



```
[29] ".01 .01 .1"
[30] ""
[31] ""
[32] ""
[33] ""
[34] ""
[35] ""
[36] ""
[37] ""
[38] "$SIGMA 0.1 0.1"
[39] ""
[40] ""
[41] ""
[42] ""
[43] "$ESTIMATION MAXEVAL=9999 PRINT=5 NOABORT METHOD=1 INTER MSFO=./1005.msf"
[44] "$COV PRINT=E"
[45] "$TABLE NOPRINT FILE=./1005.tab ONEHEADER ID AMT TIME EVID PRED IPRE CWRES"
[46] "$TABLE NOPRINT FILE=./1005par.tab ONEHEADER ID TIME CL Q V2 V3 KA ETA1 ETA2 ETA3"
[47] ""
[48] ""
[49] ""
[50] ""
[51] ""
[52] ""
[53] ""
[54] ""
[55] ""
[56] ""
[57] ""
[58] ""
[59] ""
[60] ""
[61] ""
[62] ""
```



```
[63] ""
```

Fix records of interest.

Listing 12:

```
> ctl$prob
[1] "1005 phase1 2 CMT like 1004 but diff. initial on V3"
                                                       Listing 13:
> ctl$prob <- sub('1005','1105',ctl$prob)</pre>
> names(ctl)
 [1] "prob"
                   "input"
                                "data"
                                              "subroutine" "pk"
[6] "error"
                   "theta"
                                "omega"
                                              "sigma"
                                                           "estimation"
[11] "cov"
                   "table"
                                "table"
                                                       Listing 14:
> names(ctl)[names(ctl)=='theta'] <- 'msfi'</pre>
> ctl$msfi <- '=../1005/1005.msf'
> ctl$omega <- NULL
> ctl$sigma <- NULL
> names(ctl)[names(ctl)=='estimation'] <- 'simulation'</pre>
> ctl$simulation <- 'ONLYSIM (1968) SUBPROBLEMS=500'
> ctl$cov <- NULL
> ctl$table <- NULL
> ctl$table <- NULL
> ctl$table <- 'DV NOHEADER NOPRINT FILE=./1105.tab FORWARD NOAPPEND'
> write.nmctl(ctl,'../nonmem/ctl/1105.ctl')
```



3.2 Run the simulation.

This run makes the predictions (simulations).

Listing 15:

```
> NONR72(
+ run=1105,
+ #command=command,
+ project='../nonmem',
+ grid=FALSE,
+ nice=TRUE,
+ diag=FALSE,
+ streams='../nonmem/ctl'
+ )
```

3.3 Recover and format the original dataset.

Now we fetch the results and integrate them with the other data.

Listing 16:

```
> phase1 <- read.csv('../data/derived/phase1.csv',na.strings='.')</pre>
> head(phase1)
    C ID TIME SEQ EVID AMT
                            DV SUBJ HOUR TAFD TAD LDOS MDV HEIGHT WEIGHT
    C 1 0.00 0
                      NA 0.000
                                                                74.2
                   0
                                 1 0.00 0.00
                                            NA NA
                                                           174
2 <NA> 1 0.00 1 1 1000
                           NA
                               1 0.00 0.00 0.00 1000 1
                                                           174
                                                                74.2
3 <NA> 1 0.25 0 0
                      NA 0.363
                               1 0.25 0.25 0.25 1000 0
                                                           174
                                                               74.2
4 <NA> 1 0.50 0 0 NA 0.914 1 0.50 0.50 0.50 1000 0
                                                          174 74.2
5 <NA> 1 1.00 0 NA 1.120
                                1 1.00 1.00 1.00 1000 0
                                                          174 74.2
6 <NA> 1 2.00 0 NA 2.280
                                1 2.00 2.00 2.00 1000 0
                                                          174 74.2
 SEX AGE DOSE FED SMK DS CRCN predose zerodv
```



```
1 0 29.1 1000 1
                     0 0 83.5
                                            0
2 0 29.1 1000 1
                     0 0 83.5
                                     0
                                            0
3 0 29.1 1000 1 0 0 83.5
                                            0
4 0 29.1 1000 1 0 0 83.5
                                     0
                                           0
5 0 29.1 1000
                     0 0 83.5
                                     0
                                            0
6 0 29.1 1000 1 0 0 83.5
                                                    Listing 17:
> phase1 <- phase1[is.na(phase1$C),c('SUBJ','TIME','DV')]</pre>
> records <- nrow(phase1)</pre>
> records
[1] 550
                                                    Listing 18:
> phase1 <- phase1[rep(1:records,500),]</pre>
> nrow(phase1)
[1] 275000
                                                    Listing 19:
> phase1$SIM <- rep(1:500,each=records)</pre>
> #head(phase1,300)
> with(phase1,DV[SIM==1 & SUBJ==12])
        NA 2.260 2.830 8.730 19.300 15.200 16.200 8.830 12.900 12.700
[11] 7.140 5.740 1.980 0.791
                                                    Listing 20:
> with(phase1,DV[SIM==2 & SUBJ==12])
        NA 2.260 2.830 8.730 19.300 15.200 16.200 8.830 12.900 12.700
[11] 7.140 5.740 1.980 0.791
```



3.4 Recover and format the simulation results.

```
Listing 21:

> pred <- scan('../nonmem/1105/1105.tab')
> nrow(phase1)

[1] 275000

Listing 22:

> length(pred)

[1] 275000
```

3.5 Combine the original data and the simulation data.

```
Listing 23:
```



```
SUBJ TIME DV SIM PRED
3 1 0.25 0.363 1 0.72542
4 1 0.50 0.914 1 1.38320
5 1 1.00 1.120 1 2.06720
6 1 2.00 2.280 1 3.48570
7 1 3.00 1.630 1 5.44600
8 1 4.00 2.040 1 2.99140
```

3.6 Plot predictive checks.

3.6.1 Aggregate data within subject.

Since subjects may contribute differing numbers of observations, it may be useful to look at predictions from a subject-centric perspective. Therefore, we wish to calculate summary statistics for each subject, (observed and predicted) and then make obspred comparisons therewith.

Listing 25:

> head(phase1) SUBJ TIME DV SIM PRED 1 0.25 0.363 1 0.72542 1 0.50 0.914 1 1.38320 1 1.00 1.120 1 2.06720 1 2.00 2.280 1 3.48570 1 5.44600 1 3.00 1.630 1 4.00 2.040 1 2.99140 Listing 26: > subject <- melt(phase1, measure.var=c('DV', 'PRED'))</pre> > head(subject) SUBJ TIME SIM variable value 1 1 0.25 1 DV 0.363



```
2 1 0.50 1 DV 0.914
3 1 1.00 1 DV 1.120
4 1 2.00 1 DV 2.280
5 1 3.00 1 DV 1.630
6 1 4.00 1 DV 2.040
```

We are going to aggregate each subject's DV and PRED values using cast(). cast() likes an aggregation function that returns a list. We write one that grabs min med max for each subject, sim, and variable.

Listing 27:

```
> metrics <- function(x)list(min=min(x), med=median(x), max=max(x))</pre>
```

Now we cast, ignoring time.

Listing 28:

Note that regardless of SIM, DV (observed) is constant.

DV 0.363000 1.6100 3.0900 PRED -0.022407 4.8896 12.3770

Now we melt the metrics.

Listing 29:

```
> metr <- melt(subject, measure.var=c('min', 'med', 'max'), variable_name='metric')
> head(metr)
```



```
SUBJ SIM variable metric
                              value
                DV
                      min 0.363000
              PRED
                      min 0.725420
    1 2
3
              DV
                      min 0.363000
              PRED
                      min - 0.085238
              DV
                      min 0.363000
              PRED
                      min - 0.022407
                                                    Listing 30:
> metr$value <- reapply(
       metr$value,
       INDEX=metr[,c('SIM','variable','metric')],
       FUN=sort,
       na.last=FALSE
> metr <- data.frame(cast(metr))</pre>
> head(metr)
 SUBJ SIM metric
                    DV
                           PRED
             min 0.139 - 0.61537
             med 1.025 1.25865
3
           max 2.530 2.17620
   1 2 min 0.139 -0.35196
    1 2 med 1.025 1.20926
           max 2.530 2.42390
                                                    Listing 31:
> nrow(metr)
[1] 60000
                                                    Listing 32:
> metr <- metr[!is.na(metr$DV),] #maybe no NA
```

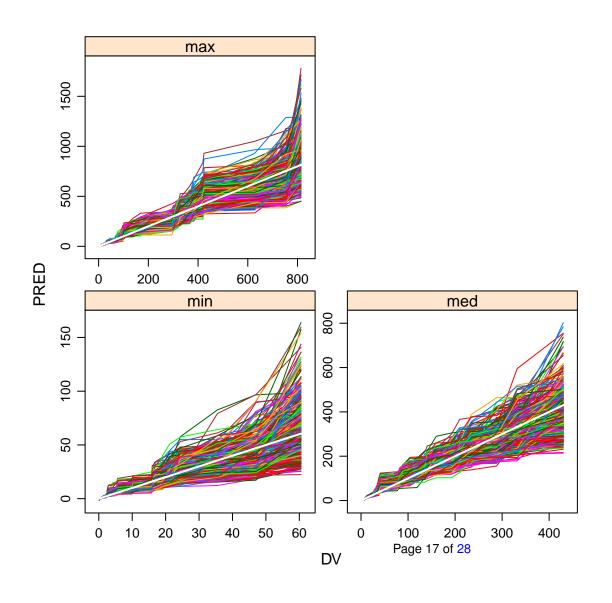


> nrow(metr)

[1] 60000

We plot using lattice.

Listing 33:





For detail, we show one endpoint, tossing the outer 5 percent of values, and indicating quartiles.

Listing 34:

```
> med <- metr[metr$metric=='med',]</pre>
> med$metric <- NULL
> head (med)
  SUBJ SIM
              DV
                   PRED
   1 1.025 1.25865
         2 1.025 1.20926
   1 3 1.025 1.57990
11
   1 4 1.025 0.88489
14
   1 5 1.025 1.65875
17
   1 6 1.025 0.95005
                                                  Listing 35:
> trim <- inner(med, id.var=c('SIM'), measure.var=c('PRED', 'DV'))</pre>
> head(trim)
 SIM DV PRED
1 1 NA
         NA
  2 NA
          NA
3
   3 NA NA
   4 NA NA
  5 NA NA
6 6 NA NA
                                                  Listing 36:
```

> nrow(trim)

[1] 20000



Listing 37:

```
> trim <- trim[!is.na(trim$DV),]
> nrow(trim)
```

[1] 19000

Listing 38:

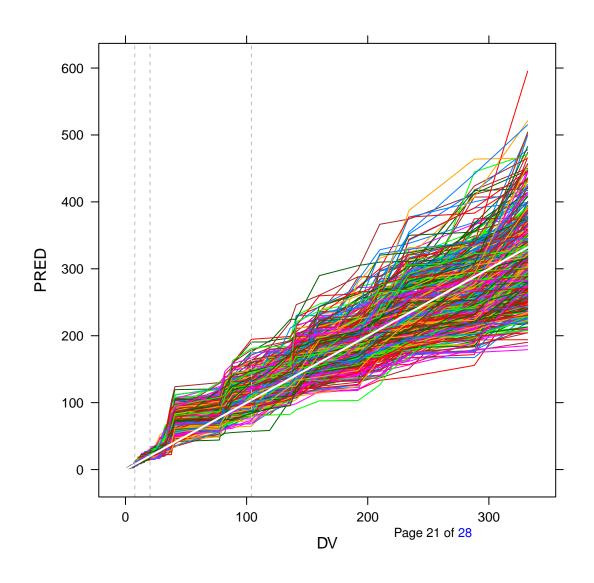
> head(trim)

```
SIM DV PRED
501 1 1.13 2.05880
502 2 1.13 2.00535
503 3 1.13 1.65480
504 4 1.13 1.06910
505 5 1.13 2.05960
506 6 1.13 0.98589
```

Listing 39:



```
+ )
```





6 6

DV 1.13

We also show densityplots of predictions at those quartiles.

```
Listing 40:
> head(trim)
               PRED
   SIM DV
501 1 1.13 2.05880
502 2 1.13 2.00535
503 3 1.13 1.65480
504 4 1.13 1.06910
505 5 1.13 2.05960
506 6 1.13 0.98589
                                                    Listing 41:
> quantile(trim$DV)
         25%
                50%
                       75% 100%
 1.13
       7.69 20.25 104.00 332.00
                                                    Listing 42:
> molt <- melt(trim, id.var='SIM')</pre>
> head(molt)
 SIM variable value
           DV 1.13
2
   2
           DV 1.13
3
   3
           DV 1.13
4
   4
           DV 1.13
           DV 1.13
```



Listing 43:

```
> quart <- data.frame(cast(molt,SIM+variable ~ .,fun=quantile,probs=c(0.25,0.5,0.75)))</pre>
> head(quart)
  SIM variable
                   X25.
                            X50.
                                      X75.
            DV 7.95000 20.25000 100.10000
2
          PRED 11.92825 22.16750 103.96500
3
          DV 7.95000 20.25000 100.10000
4
          PRED 7.23495 20.27050 105.20875
5
         DV 7.95000 20.25000 100.10000
6
          PRED 7.82690 14.50425 98.27575
                                                      Listing 44:
> molt <- melt(quart,id.var='variable',measure.var=c('X25.','X50.','X75.'),variable_name='quartile')</pre>
> head(molt)
  variable quartile
                       value
               X25. 7.95000
        DV
2
      PRED
               X25. 11.92825
3
      DV
               X25. 7.95000
4
               X25. 7.23495
      PRED
5
               X25. 7.95000
      DV
               X25. 7.82690
      PRED
                                                      Listing 45:
> levels(molt$quartile)
[1] "X25." "X50." "X75."
                                                      Listing 46:
> levels(molt$quartile) <- c('first quartile','second quartile','third quartile')</pre>
> head(molt)
```



```
variable quartile value

1 DV first quartile 7.95000

2 PRED first quartile 11.92825

3 DV first quartile 7.95000

4 PRED first quartile 7.23495

5 DV first quartile 7.95000

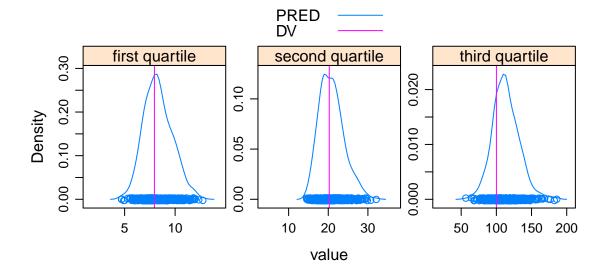
6 PRED first quartile 7.82690
```

Listing 47:

> levels(molt\$variable)

```
[1] "DV" "PRED"
```

Listing 48:





4 Bootstrap Estimates of Parameter Uncertainty

4.1 Create directories.

```
Listing 49:

> getwd()

[1] "/data/metrumrg/inst/example/project/script"

Listing 50:

> dir.create('../nonmem/1005.boot')
 > dir.create('../nonmem/1005.boot/data')
 > dir.create('../nonmem/1005.boot/ctl')
```

4.2 Create replicate control streams.

Listing 51:



```
+ ),
+ fixed=TRUE,
+ out='../nonmem/1005.boot/ctl',
+ suffix='.ctl'
+ )
```

4.3 Create replicate data sets by resampling original.

Listing 52:

```
> bootset <- read.csv('../data/derived/phase1.csv')
> r <- resample(
+ bootset,
+ names=1:300,
+ key='ID',
+ rekey=TRUE,
+ out='../nonmem/1005.boot/data',
+ stratify='SEX'
+ )</pre>
```

4.4 Run bootstrap models.

Listing 53:

```
> NONR72(
+ run=1:300,
+ #command=command,
+ project='../nonmem/1005.boot/',
+ boot=TRUE,
+ nice=TRUE,
+ grid=TRUE,
+ #concurrent=TRUE,
+ streams='../nonmem/1005.boot/ctl'
+ )
```



Installing SIGCHLD signal handler...Done.

Listing 54:

5 File Disposition

Predictive checks and bootstraps make huge files that need not be retained.

Listing 55:

```
> unlink('../nonmem/1105',recursive=TRUE)
> unlink('../nonmem/1005.boot',recursive=TRUE)
```