

## Modeling

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## 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)  
  
metrumrg 5.4  
enter "?metrumrg" for help
```

Listing 2:

```
> 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 3:

```
> 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/*/diagnostics.pdf',
+     streams='..../nonmem/ctl',
+     checksum=FALSE
+ )
```

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

Listing 4:

```
> log <- rlog(1001:1005,'..../nonmem',tool='nm7')
> 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 5:

```
> 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 phasel 2 CMT like 1004 but diff. initial on V3
249                               0
250                         ../../data/derived/phasel1.csv

```

**Listing 6:**

```

> sapply(log,class)

      tool      run   parameter      moment      value
"character" "integer" "character" "character" "character"

```

**Listing 7:**

```

> 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 8:**

```

> log <- log[log$parameter %in% c('ofv','prob','cov','min'),]
> 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                        0
39                         1001 phase1 1CMT
40                        0
42                         2525.96526753388
112                        1
113                         1002 phase1 2 CMT
114                        1
116                         2569.89393760215
153                        1
154 1003 phase1 2 CMT like 1002 but no eta on Q/v3 and no + err
155                        0

```

```

157                      2570.45022637547
194                      0
195      1004 phase1 2 CMT like 1003 but better bounds
196                      0
198                      2405.91625845151
247                      0
248      1005 phase1 2 CMT like 1004 but diff. initial on V3
249                      0

```

**Listing 9:**

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

```
[1] TRUE
```

**Listing 10:**

```

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

```

**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 11:

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

Strip comments and view.

Listing 12:

```
> ctl[] <- lapply(ctl,function(rec)sub(' *.*',' ',rec))  
> 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 13:

```
> ctl$prob
[1] "1005 phasel 2 CMT like 1004 but diff. initial on V3"
```

Listing 14:

```
> ctl$prob <- sub('1005','1105',ctl$prob)
> names(ctl)
[1] "prob"      "input"      "data"       "subroutine" "pk"
[6] "error"     "theta"      "omega"     "sigma"      "estimation"
[11] "cov"       "table"      "table"
```

Listing 15:

```
> names(ctl)[names(ctl)=="theta"] <- 'msfi'
> ctl$msfi <- '../1005/1005.msf'
> ctl$omega <- NULL
```

```
> ctl$sigma <- NULL
> names(ctl)[names(ctl)=='estimation'] <- 'simulation'
> ctl$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 16:

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

### 3.3 Recover and format the original dataset.

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

**Listing 17:**

```
> phase1 <- read.csv('../data/derived/phase1.csv',na.strings='.')
> head(phase1)
```

	C	ID	TIME	SEQ	EVID	AMT	DV	SUBJ	HOUR	TAFD	TAD	LDOS	MDV	HEIGHT	WEIGHT
1	C	1	0.00	0	0	NA	0.000	1	0.00	0.00	NA	NA	0	174	74.2
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	0	NA	1.120	1	1.00	1.00	1.00	1000	0	174	74.2
6	<NA>	1	2.00	0	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	1							
2	0	29.1	1000	1	0	0	83.5	0							
3	0	29.1	1000	1	0	0	83.5	0							
4	0	29.1	1000	1	0	0	83.5	0							
5	0	29.1	1000	1	0	0	83.5	0							
6	0	29.1	1000	1	0	0	83.5	0							

**Listing 18:**

```
> phase1 <- phase1[is.na(phase1$C),c('SUBJ','TIME','DV')]
> records <- nrow(phase1)
> records
```

```
[1] 550
```

**Listing 19:**

```
> phase1 <- phase1[rep(1:records,500),]
> nrow(phase1)
```

```
[1] 275000
```

Listing 20:

```
> phase1$SIM <- rep(1:500,each=records)
> #head(phase1,300)
> with(phase1,DV[SIM==1 & SUBJ==12])

[1]      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 21:

```
> with(phase1,DV[SIM==2 & SUBJ==12])

[1]      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 22:

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

[1] 275000
```

Listing 23:

```
> length(pred)

[1] 275000
```

### 3.5 Combine the original data and the simulation data.

Listing 24:

```
> phase1$PRED <- pred
> head(phase1)
```

SUBJ	TIME	DV	SIM	PRED
2	1 0.00	NA	1	0.00000
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

Listing 25:

```
> phase1 <- phase1[!is.na(phase1$DV),]
> head(phase1)
```

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 26:

```
> head(phase1)
```

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

Listing 27:

```
> subject <- melt(phasel1,measure.var=c('DV','PRED'))
> 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 28:

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

Now we cast, ignoring time.

Listing 29:

```
> subject <- data.frame(cast(subject, SUBJ + SIM + variable ~ .,fun=metrics))
> head(subject)
```

SUBJ	SIM	variable		min	med	max
1	1	1	DV	0.363000	1.6100	3.0900
2	1	1	PRED	0.725420	3.4795	5.4460
3	1	2	DV	0.363000	1.6100	3.0900
4	1	2	PRED	-0.085238	2.2941	4.6468
5	1	3	DV	0.363000	1.6100	3.0900
6	1	3	PRED	-0.022407	4.8896	12.3770

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

Now we melt the metrics.

**Listing 30:**

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

SUBJ	SIM	variable	metric	value
1	1	1	DV	min 0.363000
2	1	1	PRED	min 0.725420
3	1	2	DV	min 0.363000
4	1	2	PRED	min -0.085238
5	1	3	DV	min 0.363000
6	1	3	PRED	min -0.022407

**Listing 31:**

```
> metr$value <- reapply(
+   metr$value,
+   INDEX=metr[,c('SIM','variable','metric')],
+   FUN=sort,
+   na.last=FALSE
+ )
> metr <- data.frame(cast(metr))
> head(metr)
```

SUBJ	SIM	metric	DV	PRED
1	1	1	min	0.139 -0.61537
2	1	1	med	1.025 1.25865
3	1	1	max	2.530 2.17620
4	1	2	min	0.139 -0.35196
5	1	2	med	1.025 1.20926
6	1	2	max	2.530 2.42390

Listing 32:

```
> nrow(metr)
```

```
[1] 60000
```

Listing 33:

```
> metr <- metr[!is.na(metr$DV), ]#maybe no NA
> nrow(metr)
```

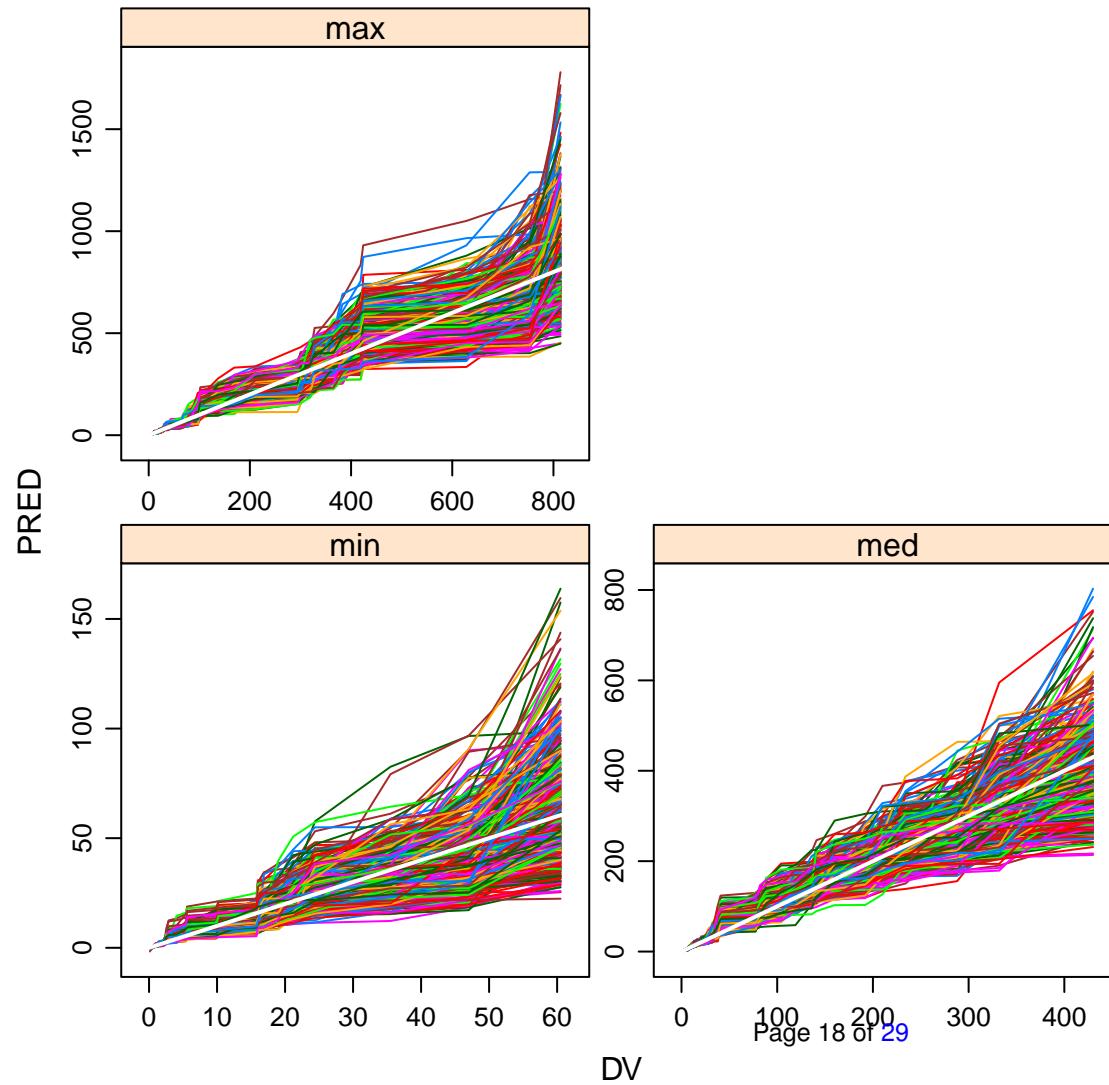
```
[1] 60000
```

We plot using lattice.

Listing 34:

```
> print(
+     xyplot(
+         PRED ~ DV|metric,
+         metr,
+         groups=SIM,
+         scales=list(relation='free'),
+         type='l',
+         panel=function(...){
+             panel.superpose(...)
+             panel.abline(0,1,col='white',lwd=2)
+         }
+     )
+ )
```

+ }  
+ )  
+ )



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

Listing 35:

```
> med <- metr[metr$metric=='med', ]  
> med$metric <- NULL  
> head(med)
```

	SUBJ	SIM	DV	PRED
2	1	1	1.025	1.25865
5	1	2	1.025	1.20926
8	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 36:

```
> trim <- inner(med, id.var=c('SIM'), measure.var=c('PRED', 'DV'))  
> head(trim)
```

	SIM	DV	PRED
1	1	NA	NA
2	2	NA	NA
3	3	NA	NA
4	4	NA	NA
5	5	NA	NA
6	6	NA	NA

Listing 37:

```
> nrow(trim)  
  
[1] 20000
```

Listing 38:

```
> trim <- trim[!is.na(trim$DV),]  
> nrow(trim)  
  
[1] 19000
```

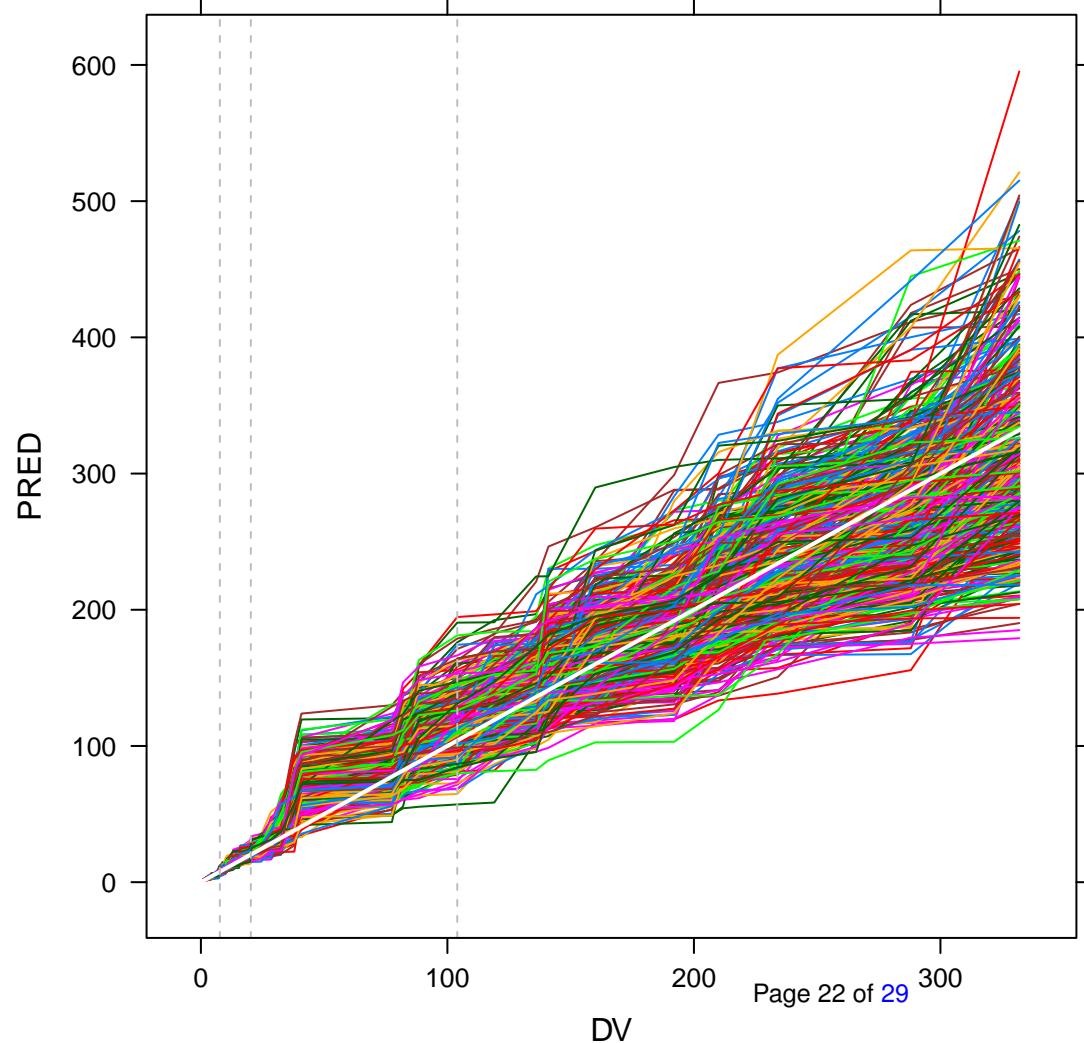
Listing 39:

```
> 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 40:

```
> print(  
+       xyplot(  
+           PRED ~ DV,  
+           trim,  
+           groups=SIM,  
+           type='l',  
+           panel=function(x,y,...){  
+               panel.xyplot(x=x,y=y,...)  
+               panel.abline(0,1,col='white',lwd=2)  
+               panel.abline(  
+                   v=quantile(x,probs=c(0.25,0.5,0.75)),  
+                   col='grey',  
+                   lty=2  
+               )
```

```
+           }  
+           )  
+   )
```



We also show densityplots of predictions at those quartiles.

Listing 41:

```
> 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 42:

```
> quantile(trim$DV)
```

	0%	25%	50%	75%	100%
	1.13	7.69	20.25	104.00	332.00

Listing 43:

```
> molt <- melt(trim, id.var='SIM')
> head(molt)
```

	SIM	variable	value
1	1	DV	1.13
2	2	DV	1.13
3	3	DV	1.13
4	4	DV	1.13
5	5	DV	1.13
6	6	DV	1.13

**Listing 44:**

```
> quart <- data.frame(cast(molt,SIM+variable ~ .,fun=quantile,probs=c(0.25,0.5,0.75)))
> head(quart)
```

	SIM	variable	X25.	X50.	X75.
1	1	DV	7.95000	20.25000	100.10000
2	1	PRED	11.92825	22.16750	103.96500
3	2	DV	7.95000	20.25000	100.10000
4	2	PRED	7.23495	20.27050	105.20875
5	3	DV	7.95000	20.25000	100.10000
6	3	PRED	7.82690	14.50425	98.27575

**Listing 45:**

```
> molt <- melt(quart,id.var='variable',measure.var=c('X25.','X50.','X75.'),variable_name='quartile')
> head(molt)
```

	variable	quartile	value
1	DV	X25.	7.95000
2	PRED	X25.	11.92825
3	DV	X25.	7.95000
4	PRED	X25.	7.23495
5	DV	X25.	7.95000
6	PRED	X25.	7.82690

**Listing 46:**

```
> levels(molt$quartile)
[1] "X25." "X50." "X75."
```

**Listing 47:**

```
> levels(molt$quartile) <- c('first quartile','second quartile','third quartile')
> head(molt)
```

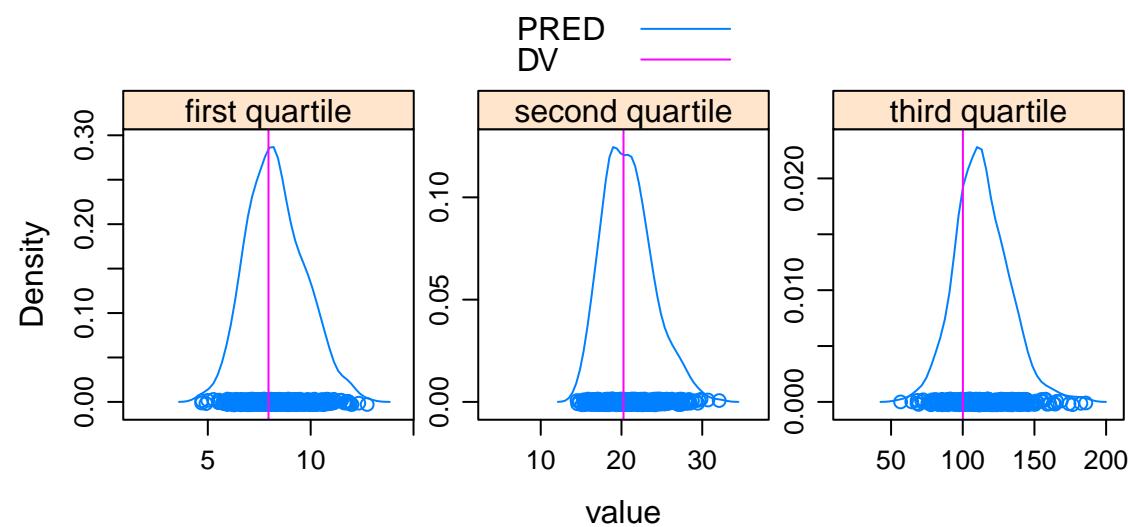
```
variable      quartile    value
1      DV first quartile 7.95000
2      PRED first quartile 11.92825
3      DV first quartile 7.95000
4      PRED first quartile 7.23495
5      DV first quartile 7.95000
6      PRED first quartile 7.82690
```

Listing 48:

```
> levels(molt$variable)
[1] "DV"     "PRED"
```

Listing 49:

```
> molt$variable <- factor(molt$variable,levels=c('PRED','DV'))
> print(
+   densityplot(
+     ~ value|quartile,
+     molt,
+     groups=variable,
+     layout=c(3,1),
+     scales=list(relation='free'),
+     aspect=1,
+     panel=panel.superpose,
+     panel.groups=function(x,...,group.number){
+       if(group.number==1)panel.densityplot(x,...)
+       if(group.number==2)panel.abline(v=unique(x),...)
+     },
+     auto.key=TRUE
+   )
+ )
```



## 4 Bootstrap Estimates of Parameter Uncertainty

### 4.1 Create directories.

Listing 50:

```
> getwd()  
  
[1] "/data/metsvn/wiki/inst/sample/script"
```

Listing 51:

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

### 4.2 Create replicate control streams.

Listing 52:

```
> t <- metaSub(  
+   clear(readLines('../nonmem/ctl/1005.ctl'),';.+',fixed=FALSE),  
+   names=1:300,  
+   pattern=c(  
+     '1005',  
+     '../..../data/derived/phasel.csv',  
+     '$COV',  
+     '$TABLE'  
+   ),  
+   replacement=c(  
+     '*',  
+     '../data/*.csv',  
+     ';$COV',  
+     ';$TABLE'
```

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

#### 4.3 Create replicate data sets by resampling original.

Listing 53:

```
> bootset <- read.csv('../data/derived/phasel1.csv')
> r <- resample(
+   bootset,
+   names=1:300,
+   key='ID',
+   rekey=TRUE,
+   out='../../nonmem/1005.boot/data',
+   stratify='SEX'
+ )
```

#### 4.4 Run bootstrap models.

Listing 54:

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

```
+ )  
Installing SIGCHLD signal handler...Done.
```

Listing 55:

```
> nms <- paste('..../nonmem/1005.boot/',1:300,'.boot/',1:300,'.log.xml',sep='')
```

```
> while(! (all(file.exists(nms)))) Sys.sleep(10)
```

```
> boot <- rlog(
```

```
+     run=1:300,
```

```
+     project='..../nonmem/1005.boot',
```

```
+     boot=TRUE,
```

```
+     append=FALSE,
```

```
+     tool='nm7'
```

```
+ )
```

```
> write.csv(boot, '..../nonmem/1005.boot/log.csv')
```