

Mlfuns Sample Script

Phase I 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:

```
> getwd()

[1] "/Users/timb/code/inst/sample/script"
```

Listing 2:

```
> library(MIfuns, lib.loc=" ~ /Rlibs")

MIfuns 4.3.3
```

Listing 3:

```
> command <- '/common/NONMEM/nm7_osx1/test/nm7_osx1.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 4:

```
> NONR(
+   run=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
+ )
> getwd()
```

```
[1] "/Users/timb/code/inst/sample/script"
```

Covariance succeeded on model 1005.

3 Predictive Check

3.1 Create a simulation control stream.

Convert control stream to R object.

Listing 5:

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

Strip comments and view.

Listing 6:

```
> 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*EXP(ERR(1)) "
[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 0.09 0.09 0.09"
[27] ""
[28] ""
```

```
[29] ""
[30] ""
[31] ""
[32] "$SIGMA 0.09"
[33] ""
[34] ""
[35] ""
[36] "$ESTIMATION MAXEVAL=9999 PRINT=5 NOABORT METHOD=1 INTER MSFO=./1005.msf"
[37] "$COV PRINT=E"
[38] "$TABLE NOPRINT FILE=./1005.tab ONEHEADER ID AMT TIME EVID PRED IPRE CWRES"
[39] "$TABLE NOPRINT FILE=./1005par.tab ONEHEADER ID TIME CL Q V2 V3 KA ETA1 ETA2
    ETA3"
[40] ""
[41] ""
[42] ""
[43] ""
[44] ""
[45] ""
[46] ""
[47] ""
[48] ""
[49] ""
[50] ""
[51] ""
[52] ""
```

Fix records of interest.

Listing 7:

```
> ctl$prob
```

```
[1] "1005 phase1 2 CMT like 1004 but diff. initial on V3"
```

Listing 8:

```
> 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 9:

```
> 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 10:

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

```
[1] "/Users/timb/code/inst/sample/script"
```

3.3 Recover and format the original dataset.

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

Listing 11:

```
> 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	1						
2	0	29.1	1000	1	0	0	83.5	0	0						
3	0	29.1	1000	1	0	0	83.5	0	0						
4	0	29.1	1000	1	0	0	83.5	0	0						
5	0	29.1	1000	1	0	0	83.5	0	0						
6	0	29.1	1000	1	0	0	83.5	0	0						

Listing 12:

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

```
[1] 550
```

Listing 13:

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

```
[1] 275000
```

Listing 14:

```
> 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 15:

```
> 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 16:

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

```
[1] 275000
```

Listing 17:

```
> length(pred)
```

```
[1] 275000
```

3.5 Combine the original data and the simulation data.

Listing 18:

```
> 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 0.17932
4     1 0.50 0.914    1 0.53642
5     1 1.00 1.120    1 0.78983
6     1 2.00 2.280    1 1.84990
7     1 3.00 1.630    1 1.96530

```

Listing 19:

```

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

```

```

SUBJ TIME    DV SIM    PRED
3     1 0.25 0.363    1 0.17932
4     1 0.50 0.914    1 0.53642
5     1 1.00 1.120    1 0.78983
6     1 2.00 2.280    1 1.84990
7     1 3.00 1.630    1 1.96530
8     1 4.00 2.040    1 2.01810

```

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 20:

```

> head(phase1)

```

```

SUBJ TIME    DV SIM    PRED
3     1 0.25 0.363    1 0.17932
4     1 0.50 0.914    1 0.53642
5     1 1.00 1.120    1 0.78983
6     1 2.00 2.280    1 1.84990
7     1 3.00 1.630    1 1.96530
8     1 4.00 2.040    1 2.01810

```

Listing 21:

```

> subject <- melt(phase1,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 22:

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

Now we cast, ignoring time.

Listing 23:

```
> 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	0.179320	1.9653	5.0314
3	1	2	DV	0.363000	1.6100	3.0900
4	1	2	PRED	0.096462	3.0448	7.4728
5	1	3	DV	0.363000	1.6100	3.0900
6	1	3	PRED	0.450430	5.5284	8.7665

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

Now we melt the metrics.

Listing 24:

```
> 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	0.179320
3	1	2	DV	min	0.363000
4	1	2	PRED	min	0.096462
5	1	3	DV	min	0.363000
6	1	3	PRED	min	0.450430

Listing 25:

```
> 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	0.064213
2	1	1	med	1.025	1.943600
3	1	1	max	2.530	3.945400
4	1	2	min	0.139	0.016162
5	1	2	med	1.025	1.476300
6	1	2	max	2.530	3.463200

Listing 26:

```
> nrow(metr)
```

```
[1] 60000
```

Listing 27:

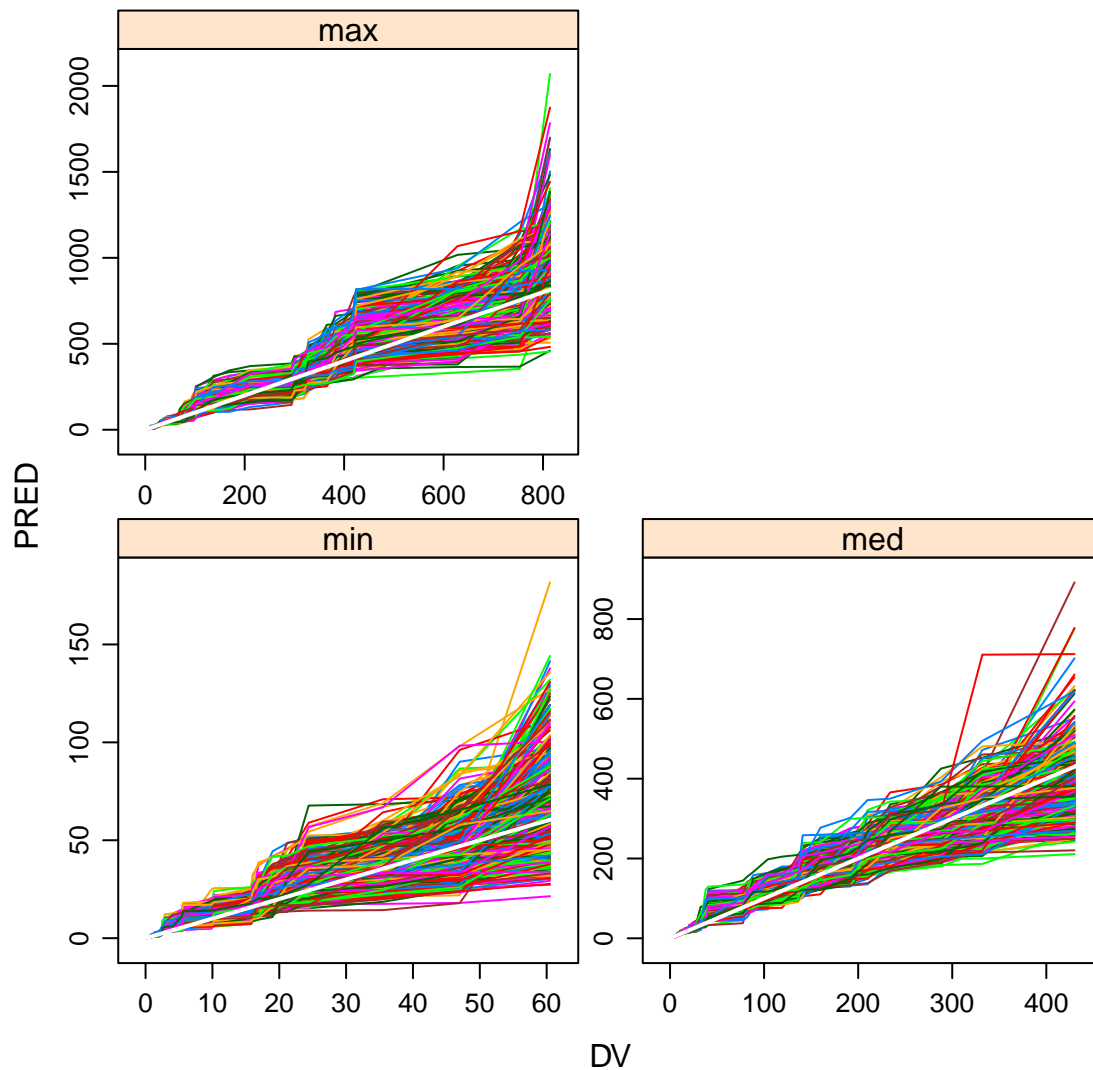
```
> metr <- metr[!is.na(metr$DV),]#maybe no NA
> nrow(metr)
```

```
[1] 60000
```

We plot using lattice.

Listing 28:

```
> 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 29:

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

	SUBJ	SIM	DV	PRED
2	1	1	1.025	1.943600
5	1	2	1.025	1.476300

```
8      1      3 1.025 1.466300
11     1      4 1.025 1.342400
14     1      5 1.025 1.362350
17     1      6 1.025 0.625815
```

Listing 30:

```
> 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 31:

```
> nrow(trim)
```

```
[1] 20000
```

Listing 32:

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

```
[1] 19000
```

Listing 33:

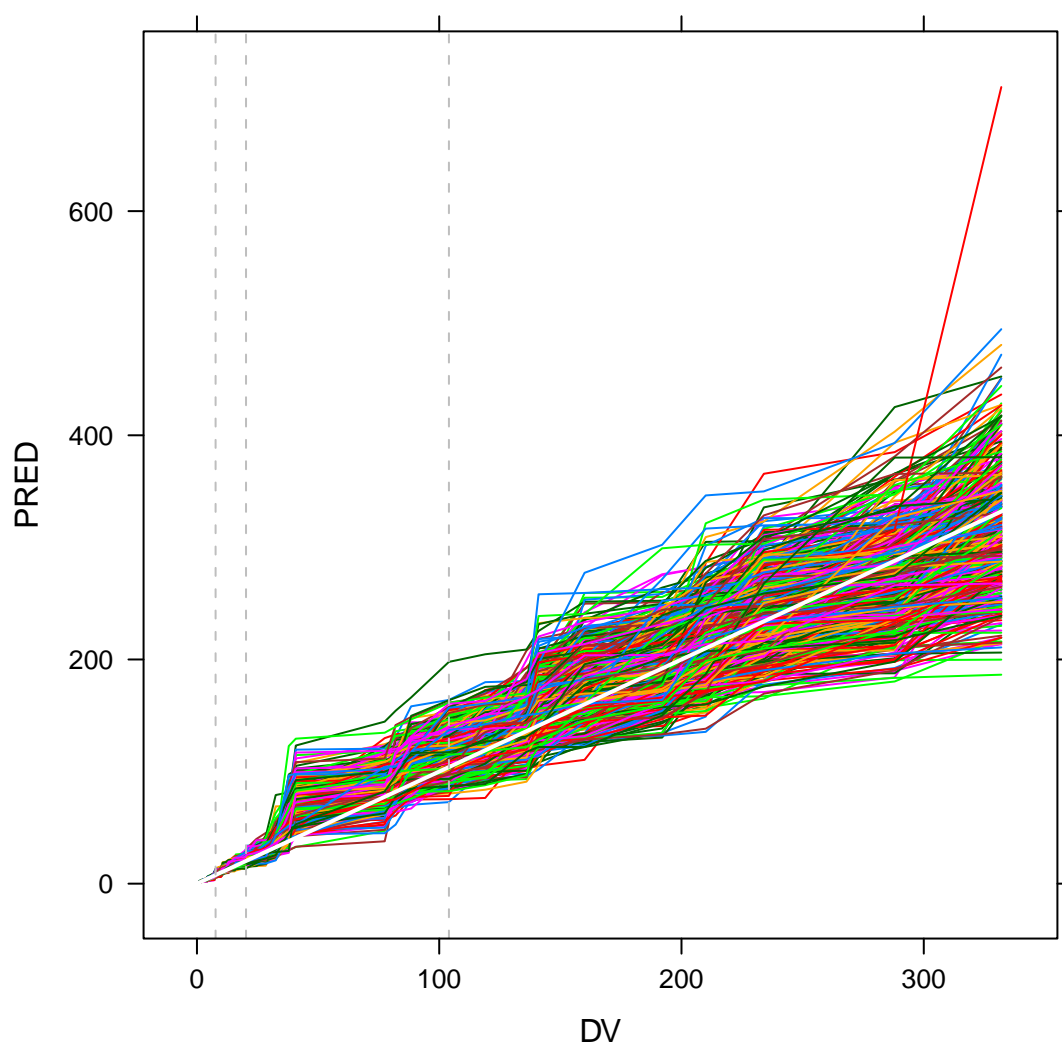
```
> head(trim)
```

```
      SIM  DV  PRED
501     1 1.13 1.9653
502     2 1.13 1.5989
503     3 1.13 1.4754
504     4 1.13 1.4074
505     5 1.13 1.3787
506     6 1.13 1.4753
```

Listing 34:

```
> 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 35:

```
> head(trim)

      SIM   DV   PRED
501    1 1.13 1.9653
502    2 1.13 1.5989
503    3 1.13 1.4754
504    4 1.13 1.4074
505    5 1.13 1.3787
506    6 1.13 1.4753
```

Listing 36:

```
> quantile(trim$DV)

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

Listing 37:

```
> 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 38:

```
> 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  7.950000 20.2500 100.10000
2     1     PRED 10.329750 22.8675  91.61825
3     2      DV  7.950000 20.2500 100.10000
4     2     PRED 10.241500 23.4225  97.26175
5     3      DV  7.950000 20.2500 100.10000
6     3     PRED  8.081437 20.0330 106.59750
```

Listing 39:

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

```

variable quartile    value
1      DV      X25.  7.950000
2     PRED      X25. 10.329750
3      DV      X25.  7.950000
4     PRED      X25. 10.241500
5      DV      X25.  7.950000
6     PRED      X25.  8.081437

```

Listing 40:

```
> levels(molt$quartile)
```

```
[1] "X25." "X50." "X75."
```

Listing 41:

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

```

variable      quartile    value
1      DV first quartile  7.950000
2     PRED first quartile 10.329750
3      DV first quartile  7.950000
4     PRED first quartile 10.241500
5      DV first quartile  7.950000
6     PRED first quartile  8.081437

```

Listing 42:

```
> levels(molt$variable)
```

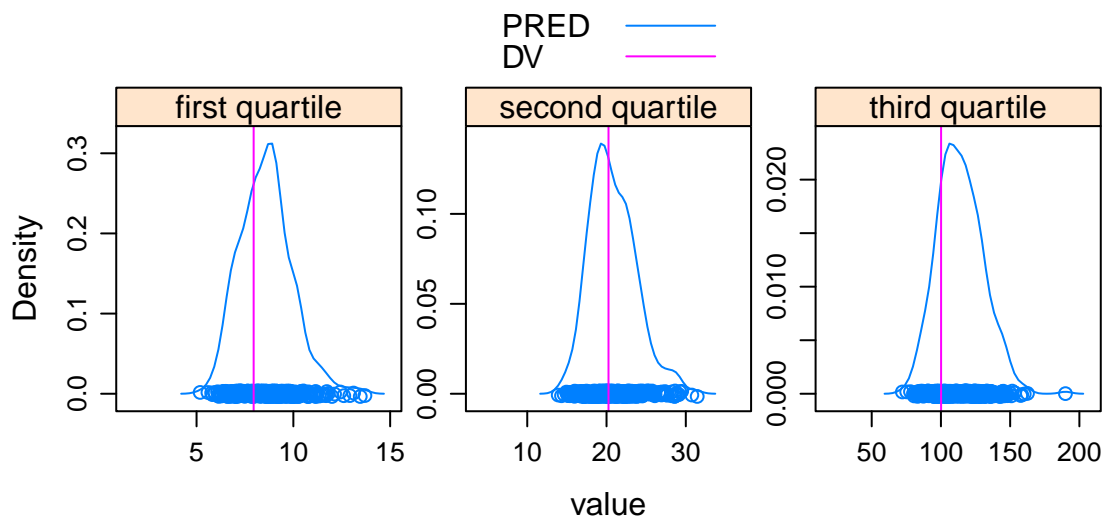
```
[1] "DV" "PRED"
```

Listing 43:

```

> 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 44:

```
> getwd()
```

```
[1] "/Users/timb/code/inst/sample/script"
```

Listing 45:

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

4.2 Create replicate control streams.

Listing 46:

```
> t <- metaSub(
+   clear(readLines('../nonmem/ctl/1005.ctl'), ';.+ ', fixed=FALSE),
+   names=1:300,
+   pattern=c(
+     '1005',
+     '../data/derived/phase1.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 47:

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

4.4 Run bootstrap models.

Listing 48:

```
> NONR (
```



```
+     run=1:300,  
+     command=command,  
+     project='../nonmem/1005.boot/',  
+     boot=TRUE,  
+     nice=TRUE,  
+     grid=FALSE,  
+     concurrent=FALSE,  
+     streams='../nonmem/1005.boot/ctl',  
+     checksum=FALSE  
+ )  
> boot <- rlog(  
+     run=1:300,  
+     project='../nonmem/1005.boot',  
+     boot=TRUE,  
+     append=FALSE,  
+     tool='nm7'  
+ )  
> write.csv(boot, '../nonmem/1005.boot/log.csv')
```