

INFUSION TESTS

kyleb

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```
Sys.setenv(RSTUDIO_PANDOC = "/usr/lib/rstudio-server/bin/pandoc")

.libPaths("/data/Rlibs")
library(mrgsolve)
library(dplyr)

##
## Attaching package: 'dplyr'
##
## The following objects are masked from 'package:stats':
##
##   filter, lag
##
## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
library(readr)
library(ggplot2)

carry <- c("cmt", "amt", "ii", "addl", "rate", "evid", "ss")

to_data_set <- function(x, id = NULL) {
  x <- as.data.frame(x)
  x <- mutate(x, C = '.', DV = '.', cmt = if_else(cmt==0, 2, cmt))
  x <- dplyr::select(x, "C", everything())
  if(is.numeric(id)) x <- mutate(x, ID = id)
  x
}

sv <- function(x, file) {
  write.csv(file = file, row.names = FALSE, quote = FALSE, x)
}

nonr <- metrumrg::NONR
run <- function(number) {
  nonr(number, project = "model", command = "/opt/NONMEM/nm73/nmqual/autolog.pl", checkrunno=FALSE)
  return(tabread(number))
}

tabread <- function(number) {
  tab <- file.path("model", number, "TAB")
  if(file.exists(tab)) return(read_table(tab, skip=1))
  stop("the run failed")
}

sim <- function(x, e,...) {
  mrgsim(x, events = e, carry.out = carry, digits = 5, ...)
}

code <- '
```

```

$SET req = ""
$PARAM CL = 1, V = 30, KA = 1.5
LAGT = 0, MODE = 0, DUR2 = 2, RAT2 = 10, BIOAV = 1

$PKMODEL cmt = "GUT CENT", depot = TRUE

$MAIN

F_CENT = BIOAV;
ALAG_CENT = LAGT;

if(MODE==1) R_CENT = RAT2;
if(MODE==2) D_CENT = DUR2;

$TABLE
capture DV = (CENT/(V/1000));
capture CP = DV;

$CAPTURE LAGT MODE DUR2 RAT2 BIOAV
,

mod <- mcode_cache("tests1", code)

## Compiling tests1 ...
## done.
mod <- update(mod, end=72)

```

Bolus

```

ev <- ev(amt = 100, ii = 12, addl = 3)
out1 <- sim(mod, ev)
data1 <- to_data_set(out1, 1)

```

Infusion

```

ev <- ev(amt = 100, ii = 12, addl = 3, rate = 100/10, cmt = 2)
out2 <- sim(mod, ev)
data2 <- to_data_set(out2, 2)

```

Infusion, lag

```

ev <- ev(amt = 100, ii = 12, addl = 3, rate = 100/10, LAGT = 5, cmt = 2)
out3 <- sim(mod, ev)
data3 <- to_data_set(out3, 3)

```

Infusion, lag, bioav

```
ev <- ev(amt = 100, ii = 12, addl=3, rate = 100/10, LAGT = 5, BIOAV = 0.412, cmt = 2)
out4 <- sim(mod,ev)
data4 <- to_data_set(out4, 4)
```

Infusion, bioav, ss

```
ev <- ev(amt = 100, ii = 12, addl=3, rate = 100/10, LAGT = 0, BIOAV = 0.412, ss = 1, cmt = 2)
out5 <- sim(mod,ev)
data5 <- to_data_set(out5, 5)
```

Infusion, bioav, ss, II < DUR

```
ev <- ev(amt = 100, ii = 6, addl = 3, rate = 100/10, BIOAV = 0.812, ss = 1, cmt = 2)
out6 <- sim(mod,ev)
data6 <- to_data_set(out6, 6)
```

Infusion, ss, II < DUR

```
ev <- ev(amt = 100, ii = 6, addl = 3, rate = 100/10, ss = 1, cmt = 2)
out6.1 <- sim(mod,ev)
data6.1 <- to_data_set(out6.1, 6.1)
```

Infusion, ss, II multiple of DUR

```
ev <- ev(amt = 100, ii = 6, addl = 3, rate = signif(100/12,5), ss = 1, cmt = 2)
out6.2 <- sim(mod,ev)
data6.2 <- to_data_set(out6.2, 6.2)
```

Infusion, bioav, ss, II == DUR

```
ev <- ev(amt = 100, ii = 10, addl=3, rate = 100/10, LAGT = 0, BIOAV = 0.412, ss = 1, cmt = 2)
out7 <- sim(mod,ev)
data7 <- to_data_set(out7, 7)
```

Infusion,, ss, II == DUR

```
ev <- ev(amt = 100, ii = 10, addl=3, rate = 100/10, LAGT = 0, ss = 1, cmt = 2)
out7.1 <- sim(mod,ev)
data7.1 <- to_data_set(out7.1, 7.1)
```

Bolus, bioav, ss

```
ev <- ev(amt = 100, ii = 12, addl=3, LAGT = 0, BIOAV = 0.412, ss = 1)
out8 <- sim(mod, ev)
data8 <- to_data_set(out8, 8)
```

Bolus, lag, bioav

```
ev <- ev(amt = 100, ii = 12, addl=3, LAGT = 5, BIOAV = 0.412)
out9 <- sim(mod, ev)
data9 <- to_data_set(out9, 9)
```

Infusion / bolus

```
ev <- ev(amt = 100, rate = 10) + ev(time = 12, amt = 50)
out10 <- sim(mod, ev)
data10 <- to_data_set(out10, 10)
```

Infusion (D_) lag, BIOAV

```
ev <- ev(amt = 100, rate = -2, DUR2 = 9, MODE = 2, cmt = 2, ii = 12, addl = 5, LAGT = 5, BIOAV = 0.61)
out11 <- sim(mod, ev)
data11 <- to_data_set(out11, 11)
```

Simulate

```
sims <- list(out1, out2, out3, out4, out5, out6, out6.1, out6.2, out7, out7.1, out8, out9, out10, out11)
sims <- lapply(sims, as.data.frame)
sims <- bind_rows(sims)
data <- bind_rows(data1, data2, data3, data4, data5, data6, data6.1, data6.2, data7, data7.1, data8, data9, data10,
  sv(data, "data/101.csv")
out <- run(101)
```

```
## Run 101 complete.
```

```
## NONR complete.
```

```
## Parsed with column specification:
```

```
## cols(
```

```
##   TIME = col_double(),
```

```
##   EVID = col_double(),
```

```
##   CP = col_double(),
```

```
##   IPRED = col_double(),
```

```
##   PRED = col_double(),
```

```
##   DV = col_double()
```

```
## )
```

Overall Summary

```
dim(out)

## [1] 1037    6

dim(sims)

## [1] 1037    16

summary(out$CP - sims$CP)

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##         0         0         0         0         0         0

data$NM <- out$CP
data$MRGSIM <- sims$CP
```

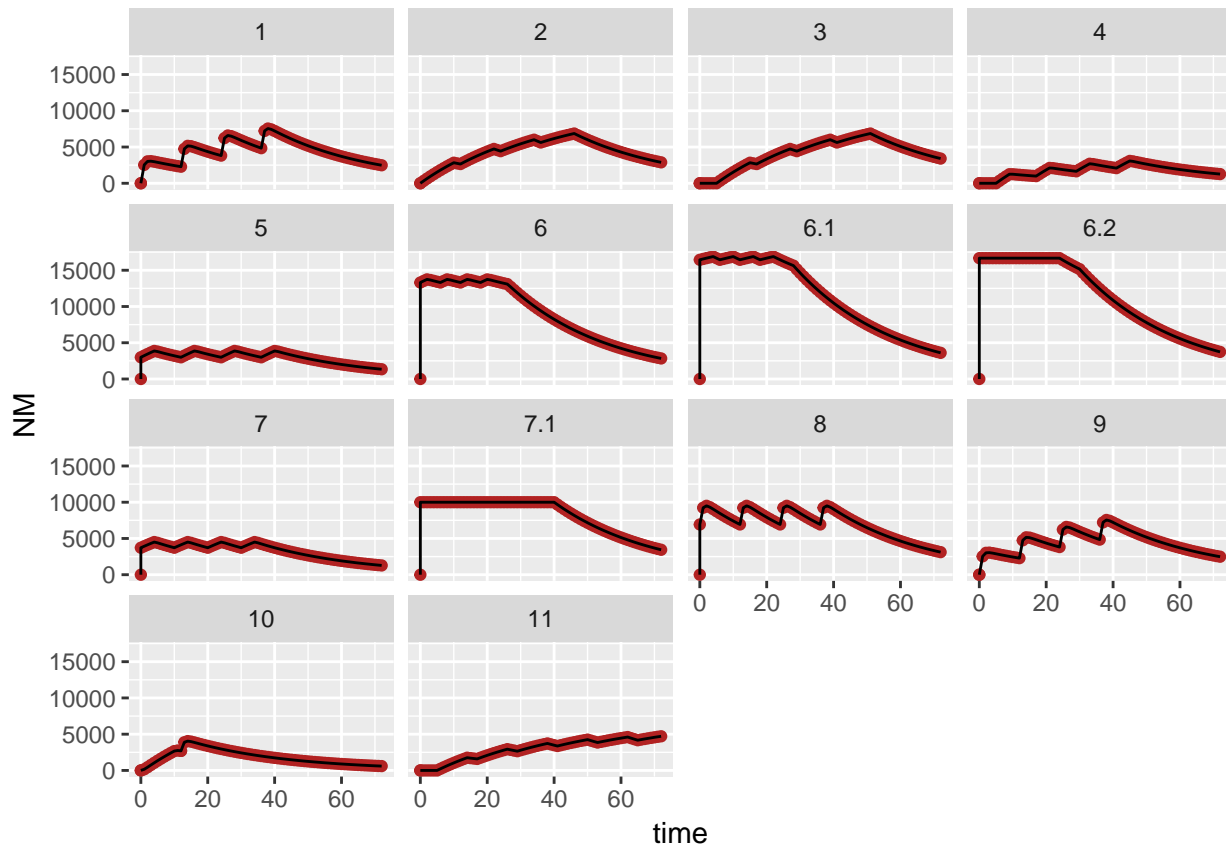
Summary by RUN

```
group_by(data,ID) %>%
  mutate(diff = NM - MRGSIM) %>%
  summarise(mean = mean(diff), max = max(diff), min = min(diff))

## # A tibble: 14 x 4
##       ID mean  max  min
##   <dbl> <dbl> <dbl> <dbl>
## 1  1.0    0    0    0
## 2  2.0    0    0    0
## 3  3.0    0    0    0
## 4  4.0    0    0    0
## 5  5.0    0    0    0
## 6  6.0    0    0    0
## 7  6.1    0    0    0
## 8  6.2    0    0    0
## 9  7.0    0    0    0
## 10 7.1    0    0    0
## 11 8.0    0    0    0
## 12 9.0    0    0    0
## 13 10.0   0    0    0
## 14 11.0   0    0    0
```

Plot

```
ggplot(data = data) +
  geom_point(aes(time,NM),color = "firebrick") +
  geom_line(aes(time,MRGSIM,group = ID)) +
  facet_wrap(~ID)
```



Control stream

```
writeLines(readLines("model/101.ct1"))
```

```
$PROB RUN# 101
```

```
$INPUT C ID TIME EVID AMT CMT SS II ADDL RATE LAGT MODE DUR2 RAT2 BIOAV DV
```

```
$DATA ../../data/101.csv IGNORE=C
```

```
$SUBROUTINES ADVAN2 TRANS2
```

```
$PK
```

```
TVCL=THETA(1)
```

```
CL=TVCL*EXP(ETA(1))
```

```
TVV2=THETA(2)
```

```
V=TVV2*EXP(ETA(2))
```

```
TVKA=THETA(3)
```

```
KA=TVKA*EXP(ETA(3))
```

```
ALAG2 = LAGT
```

```
F2 = BIOAV
```

```
IF(MODE.EQ.1) R2 = RAT2
IF(MODE.EQ.2) D2 = DUR2
```

```
$ERROR
IPRED=A(2)/(V/1000)
Y=IPRED*EXP(ERR(1))
```

```
CP = IPRED
```

```
$THETA
(1, FIX) ;; CL
(30, FIX) ;; V
(1.5, FIX) ;; KA
```

```
$OMEGA
0.0 FIX
0.0 FIX
0.0 FIX
```

```
$SIGMA
0.00 FIX
```

```
$TABLE FILE=TAB TIME EVID CP IPRED PRED DV NOPRINT ONEHEADER NOAPPEND
```

```
$SIMULATION (2674474) ONLYSIMULATION
```

```
devtools::session_info()
```

```
## Session info -----
```

```
## setting value
## version R version 3.3.3 (2017-03-06)
## system x86_64, linux-gnu
## ui X11
## language (EN)
## collate en_US.UTF-8
## tz Etc/UTC
## date 2017-09-15
```

```
## Packages -----
```

## package	* version	date
## assertthat	0.2.0	2017-04-11
## backports	1.1.0	2017-05-22
## base	* 3.3.3	2017-03-06
## bindr	0.1	2016-11-13
## bindrcpp	* 0.2	2017-06-17
## colorspace	1.3-2	2016-12-14
## datasets	* 3.3.3	2017-03-06
## devtools	1.13.2	2017-06-02
## digest	0.6.12	2017-01-27
## dplyr	* 0.7.3	2017-09-09
## evaluate	0.10.1	2017-06-24
## fork	1.2.5	2017-07-26
## ggplot2	* 2.2.1	2016-12-30

```

## glue                1.1.1      2017-06-21
## graphics            * 3.3.3      2017-03-06
## grDevices           * 3.3.3      2017-03-06
## grid                3.3.3      2017-03-06
## gtable              0.2.0      2016-02-26
## hms                 0.3         2016-11-22
## htmltools           0.3.6      2017-04-28
## knitr               1.16        2017-05-18
## labeling            0.3         2014-08-23
## lattice             0.20-35     2017-03-25
## lazyeval            0.2.0      2016-06-12
## magrittr            1.5         2014-11-22
## MASS               7.3-45      2016-04-21
## memoise             1.0.0      2016-01-29
## methods             * 3.3.3      2017-03-06
## metrumrg            5.57        2015-10-08
## mrgsolve            * 0.8.9.9001 2017-09-15
## munsell              0.4.3      2016-02-13
## pkgconfig           2.0.1      2017-03-21
## plyr                1.8.4      2016-06-08
## R6                  2.2.2      2017-06-17
## Rcpp                0.12.12     2017-07-15
## RcppArmadillo       0.7.960.1.2 2017-08-29
## readr               * 1.1.1      2017-05-16
## reshape            0.8.6        2016-10-21
## rlang               0.1.2      2017-08-09
## rmarkdown           1.6         2017-06-15
## rprojroot           1.2         2017-01-16
## scales              0.5.0      2017-08-24
## stats               * 3.3.3      2017-03-06
## stringi             1.1.5      2017-04-07
## stringr             1.2.0      2017-02-18
## tibble              1.3.4      2017-08-22
## tools               3.3.3      2017-03-06
## utils               * 3.3.3      2017-03-06
## withr               1.0.2      2016-06-20
## XML                 3.98-1.9  2017-06-19
## yaml                2.1.14     2016-11-12
## source
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## local
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.2)
## local
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## cran (@0.7.3)
## CRAN (R 3.3.3)
## local
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## local

```



```

## local
## local
## CRAN (R 3.3.2)
## CRAN (R 3.2.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.2)
## CRAN (R 3.3.3)
## CRAN (R 3.2.3)
## CRAN (R 3.3.2)
## CRAN (R 3.3.2)
## CRAN (R 3.3.2)
## local
## R-Forge (R 3.2.3)
## Github (metrumresearchgroup/mrgsolve@8cbf215)
## CRAN (R 3.3.2)
## CRAN (R 3.3.3)
## CRAN (R 3.3.2)
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## cran (@0.7.960)
## CRAN (R 3.3.3)
## CRAN (R 3.2.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## local
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## CRAN (R 3.3.3)
## local
## local
## CRAN (R 3.3.2)
## CRAN (R 3.3.3)
## CRAN (R 3.3.2)

```