



mrgsolve: Some ideas

mrgsolve Workshop

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San Diego, CA

Decide which updates to make persistent

Sometimes you want to do this:

```
mod %<>% update(end=2400)
out <- mod %>% mrgsim
```

Other times this:

```
out <- mod %>% mrgsim(end=2400)
```

Parallelize your simulation

- ▶ Most frequently we parallelize across posterior draws
 - ▶ Each posterior draw or bootstrap sample goes to a different worker
- ▶ When the data set is large, you can split it on an appropriate factor
 - ▶ Study
 - ▶ Treatment arm
 - ▶ Treatment duration
- ▶ Use `mclapply` in Mac/Unix and `doParallel` in Windows
 - ▶ Only utilizes cores on the master node
- ▶ We use a `qapply` package that will send jobs to all nodes in a cluster via Grid Engine

Save simulated output in serialized format (if needed)

```
out <- mod %>% mrgsim  
saveRDS(file="mysim.RDS", out)
```

```
out <- readRDS(file="mysim.RDS")
```

Summarize the simulation prior to returning from the worker

Helpful when memory is limited

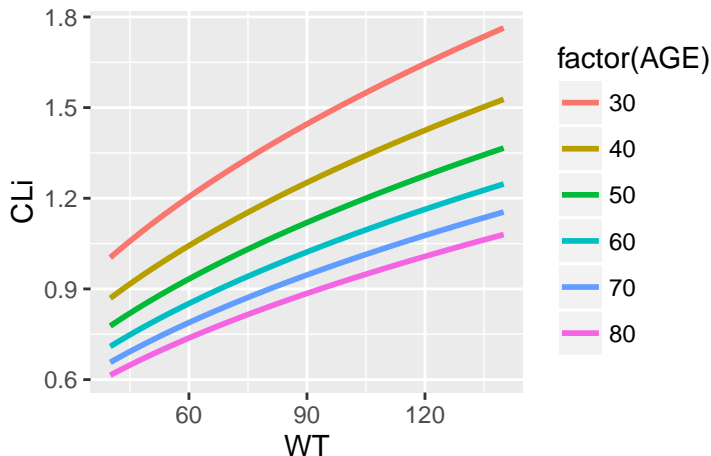
```
out <- mclapply(1:1000, function(i) {  
  
  mod %>%  
    data_set(...) %>%  
    mrgsim %>%  
    mutate(irep=i) %>%  
    filter(time > 168) %>%  
    group_by(ID,DOSE) %>%  
    summarise(Cmax=max(DV))  
  
}) %>% bind_rows
```

Take advantage of the covariate model

```
code <- '  
$PARAM TVCL=1, TVVC=20, WT=70, AGE=50  
  
$MAIN  
double CLi = TVCL*pow(WT/70,0.45)*pow(AGE/50,-0.5)*exp(ETA  
  
$CAPTURE CLi WT AGE  
  
// Other stuff here....  
'  
  
mod <- mread("covmodel", tempdir(),code, warn=FALSE)
```

CL vs WT by AGE

```
idata <- expand.idata(WT=seq(40,140),AGE=seq(30,80,10))  
out <- mod %>% idata_set(idata) %>% mrgsim(end=-1)
```



Extend mrgsolve

```
doseA <- function(x,amt,mw=1.23,ii=12,cmt=2,...) {  
  x %>% ev(amt=amt/mw/1000,cmt=cmt,ii=ii,...)  
}
```

```
mod %>%  
  doseA(45,addl=23) %>%  
  mrgsim(Req="CENT") %>% head(n=2)
```

```
. Model:  housemodel
```

```
.      ID time      CENT  
. [1,]  1    0 0.00000000  
. [2,]  1    0 0.03658537
```


Start developing the simulation model early

- ▶ Don't wait until the end of the project to translate the model to `mrgsolve` so you can simulate all of the different regimens
- ▶ When you start simulating early, you will use / re-use the model more