

PK/PD Modeling with RsNLME : : CHEAT SHEET

Basics

RsNLME is an R package to define NLME models as R native objects.

- NLME engine is used for Fitting/Simulation
- R style help available via ?method
- All methods have sensible default values

Model Creation

```
pk = pkmodel(numComp = 1,
             absorption=Intravaneous,...)
pd = emaxmodel(checkBaseline=TRUE,...)
pd = linearmodel(type = LinearAlpha,...)
pkpd = pkemaxmodel(numComp = 2,
                  absorption=Extravascular
                  isTlag = TRUE,...)
pkpd = pkindirectmodel(isClosedForm=FALSE,
                      indirectType = LimitedStimulation,...)
pkpd = pklinearmodel(parameterization= Micro,
                    linearModelType=LinearBeta, ...)
model = blankmodel(modelName, ...)
```

Input Options

```
model2 = addReset(model, low, hi)
model2 = addExtraDose(model, doseType, doses)
Dose Types : SteadyStateDose | AddIDose
```

Model covariates

```
sex=categoricalCovariate("sex",c(1,2),c("female","male"))
weight=NlmeCovariateParameter("weight",centerValue="70",
                              continuousType =CovarNumber,
                              direction=Forward)
age=NlmeCovariateParameter("age")
occ=occasionCovariate("OCC",c(1,2), c("OCC1","OCC2"),
                     direction=Forward)
model=addCovariates(model, c(sex,weight,age,occ),
                   c("V"="weight,age", "Cl"="sex,weight"))
newModel = resetCovariateEffects(model)
covariateEffect(model,"wt","Cl")=COVAR_EFF_YES
```

Customizing the model

Residual error model

```
residualEffect(model,"C")=c(errorType=Multiplicative,SD="0.16",
                             isFrozen=FALSE,isBQL=TRUE,bqlStatic=0.75,
                             doBefore="",doAfter="")
```

Error model types :

Additive | LogAdditive | Multiplicative | AdditiveMultiplicative | MixRatio | Power | Custom

```
structuralParam(model,"V") = c(style=Custom,
                                code="stparm(V=10^(tvlog10V + nlog10V))")
```

Random effects

```
initRandomEffects(model)=c(Block, FALSE,
                             "nV,nCl,nKa,nV2", "0.2, 0, 0.2, 0, 0, 0.2, 0, 0, 0.1")
initRandomEffects(model)=c(Diagonal,
                             FALSE,"nV,nCl","0.1, 0.02")
initRandomEffects(model)=c(
    Diagonal, FALSE,"nV,nCl","0.1, 0.02",
    Block, TRUE, "nCl2,nV2","0.2, 0, 0.2")
initOccasionRandomEffect(model,"Occasion") =
c(0.1,0.02,0.1)
```

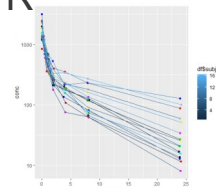
Add more components to the model

```
model2 = addCountObservation(model, observationName,
                             .....expression, structuralParameters)
model2 = addCategoricalObservation(model,
                                   observationName, offsetArray,
                                   structuralParameters)
model2 = addContinuousObservation(model,
                                   observationName,effect, hasRandomEffect)
model2 = addLLObservation(model,observationName,
                           expression,dobefore,doafter, structuralParameters,
                           isFrozen, hasRandomEffect,simulationCode)
model2 = addEventObservation(model,observationName,
                              epression,dobefore,doafter,structuralParameters,
                              isFrozen, hasRandomEffect)
model2 = addParameters(model,name,...)
model2= addExpression(model,blockName
                      ,structuralParameters, codeLine, isFrozen, override)
```

Input dataset

Visualize the data with R

```
input=read.csv("16subjects.csv")
ggplot(data=input,aes(x=time,y=conc))+
  scale_y_log10()+
  geom_point(colour=df$subject)+
  geom_line(aes(x=df$time,y=df$conc,
               group=df$subject,colour=df$subject))
```

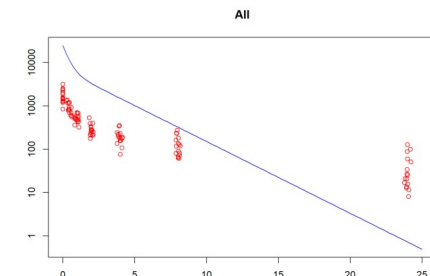
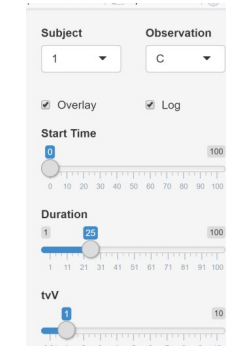


Map columns to model variables

```
initColMapping(model)=input
print(modelColumnMapping(model))
modelColumnMapping(model)=c(id="Subject",
                             CObs="Conc",A1="Amount")
```

Initial estimates for fixed effects

```
estimatesUI(model,unique(input$Subject),host)
```



```
effects=getInitialEstimates()
initFixedEffects(model) = effects
```

Platform/Engine defintion

```
method=NlmeParallelMethod(method="MULTICORE")["MP
I"]["TORQUE"]["TORQUE_MPI"]["SGE"]["SGE_MPI"]["LSF"]["LS
F_MPI"]
host = NlmeParallelHost(sharedDirectory="C:/Shared",
    parallelMethod=method,
    hostName="MPI",
    numCores=4)
```

Fitting Engines Definition

```
engineParams = NlmeEngineExtraParams(
    PARAMS_METHOD=METHOD_FOCE_ELS,
    PARAMS_NUM_ITERATIONS=1000,
    PARAMS_SAND="TRUE")

METHOD_QRPEM          METHOD_FIRST_ORDER
METHOD_IT2S_EM        METHOD_LAPLACIAN
METHOD_FOCE_LB        METHOD_NAIVE_POOLED
```

Model Fitting

```
Job = fitmodel(hostPlatform, params , model,
    runInBackground )
```

Ex.

```
job = fitmodel(host,engineParams)
status = read.csv("Overall.csv")
print(status)
```

Model Simulation

```
Job = simmodel(hostPlatform, simParams ,
    model , runInBackground )
```

Ex.

```
SimTableObs = NlmeSimTableDef("SimTableObs.csv",
    "0,1,2,4,4.9,55.1,56,57,59,60", "C, CObs", FALSE)
simParams = NlmeSimulationParams(numReplicates = 50,
    seed = 3527, simulationTables = c(SimTableObs))
job = simmodel(defaultHost,simParams,model)
```

Covariate Search

```
stepwiseSearch(hostPlatform, params,
    covariateModel, stepwiseParams, model,
    runInBackground)
```

```
shotgunSearch(hostPlatform, params,
    covariateModel, model, runInBackground)
```

Example

```
sp = NlmeStepwiseParams(0.01, 0.001, "-2LL")
job=stepwiseSearch(host, params,
    covariateModel(model), sp, model)
```

Bootstrap

```
bootstrap(hostPlatform, params, bootParams, model,
    runInBackground)
```

Ex.

```
bootParams = NlmeBootstrapParams(
    numReplicates=5, randomNumSeed=1234)
job = bootstrap(defaultHost, params,
    bootParams, model,TRUE)
```

VPC

```
vpcmodel(hostPlatform, vpcParams, model=model, ...)
```

Ex.

```
obsVars = GetObservationVariables(model@dataset)
observationParameters(obsVars[[1]])=c(xaxis=VPC_XAXIS_T,
    binningMethod=VPC_BIN_NONE,
    quantilesValues ="5,50,95") vpcParams =
    NlmeVpcParams(numReplicates=2, seed=1234,
    observationVars=obsVars)
job = vpcmodel(defaultHost ,vpcParams ,model)
```

Scenario Fitting

```
scenario1=NlmeScenario("SC0001","1")
scenario2=NlmeScenario("SC0002","1,2")
scenarios = c(scenario1,scenario2)
sortColumns=NlmeSortColumns("group,sex")
job = sortfit(defaultHost,params,
    sortColumns,scenarios,model)
```

Analyzing Results

VPC

```
library(vpc)
simData = getSimData(input,stratifyColumns="sex")
obsData = getObsData(input)
vpcdb = vpc(sim=simData, obs = obsData, vpcdb = TRUE)
plot_vpc(vpcdb,
    show = list(obs_dv = TRUE, obs_ci = FALSE),
    xlab = "Time(hours)", ylab = "Concentration",
    title = "VPC!")
```

Diagnostic plots

```
xp = xposeNlme(dir=".",modelName="Initial Model")
list_vars(xp)
doexpose(xp)
dv_vs_pred(xp)
res_vs_pred(xp,res="CWRES",type="ps")
ind_plots(xp)
eta_distrib(xp)
eta_qq(xp)
```

Bootstrap

```
out=read.csv("out.csv")
View(out)
overall=read.csv("BootOverall.csv")
View(overall)
theta=read.csv("BootTheta.csv")
View(theta)
varCovar=read.csv("BootVarCoVar.csv")
View(varCovar)
omega=read.csv("BootOmega.csv")
View(omega)
```

Covariate Search

```
overall = read.csv("Overall.csv")
View(overall)
stepwiseLines=readLines("Stepwise.txt")
View(stepwiseLines)
```
