



mrgsolve: Model Specification

mrgsolve Workshop

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San Diego, CA

Available code blocks

- ▶ \$PARAM, \$THETA, \$FIXED
- ▶ \$CMT, \$INIT, \$VCMT
- ▶ \$MAIN
- ▶ \$ODE
- ▶ \$TABLE , \$CAPTURE
- ▶ \$OMEGA , \$SIGMA
- ▶ \$GLOBAL
- ▶ \$PROB, \$SET, \$ADVAN2, \$ADVAN4, \$NMXML, \$CMTN

Assemble blocks into something like this

```
$PARAM CL=1, VC=35, VMAX=2, KM=1.2 KA=1.1

$CMT SC CENT

$MAIN
double VCi = VC*exp(ETA(2));

$OMEGA 0.5 0.5

$ODE
dxdt_SC = -KA*SC;
dxdt_CENT = KA*SC - (CL/VCi)*CENT;

$TABLE table(CP) = CENT/VCi; // forget about this
```

\$PARAM [R]

- ▶ Model “parameters” with name=value format
 - ▶ Separate by , or \n
 - ▶ Use any name except for words in `mrgsolve:::reserved()`
 - ▶ Values are evaluated by R parser
- ▶ Multiple blocks allowed
- ▶ Use name anywhere in the model
- ▶ Updatable in R but not in the model spec
 - ▶ Most often, you will want to match names in \$PARAM with the names in your input data sets

\$PARAM

TVCL=1.1, TVVC=1.2

KA = 0.7, F1 = sqrt(2/3)

- ▶ See also: \$THETA and \$FIXED
- ▶ **ACCESS** R global environment

\$CMT [txt] and \$INIT [R]

- ▶ Specify name and number of compartments in the model
- ▶ Use any name except for those listed in `mrgsolve:::reserved()`
- ▶ Use either \$CMT or \$INIT
- ▶ Maybe used to specify initial condition
 - ▶ \$CMT specify compartment names only
 - ▶ \$INIT to specify `name = <initial>`

See also: \$VCMT

ACCESS R global environment for \$INIT only

The difference between \$CMT and \$INIT

- ▶ These give identical results

```
$CMT GUT CENT RESP
```

```
$INIT GUT=0, CENT=0, RESP=0
```

- ▶ But you may wish to do this

```
$INIT GUT=0, CENT=0, RESP=50
```

Variables and macros (1/3)

- ▶ CMT_0 the initial condition for CMT
- ▶ CMT the current amount in CMT
- ▶ TIME the current data set time
- ▶ SOLVERTIME in \$ODE only ... the current ode solver time
- ▶ EVID event id
- ▶ ID subject ID
- ▶ NEWIND new individual flag
 - ▶ 0 for first record of data set
 - ▶ 1 for first record of this ID
 - ▶ 2 for subsequent of this ID

Variables and macros (2/3)

- ▶ Random effects
 - ▶ $\text{ETA}(m)$
 - ▶ $\text{EPS}(m)$
- ▶ General dosing
 - ▶ $_F(n)$ bioavailability parameter
 - ▶ $_ALAG(n)$ dose lag time
- ▶ Infusion settings
 - ▶ $_D(n)$ used to set infusion duration
 - ▶ $_R(n)$ used to set infusion rate

n (compartment number) and m (ETA or EPS number) are integers

Variables and macros (3/3)

- ▶ `_NEQ` number of equations
- ▶ `_N_CMT` the number (integer) of compartment CMT; but **only** if you used `$CMTN`
- ▶ `$ADVAN2`
 - ▶ `pred_CL`, `pred_VC`, `pred_KA`
- ▶ `$ADVAN4`
 - ▶ `pred_CL`, `pred_VC`, `pred_KA`, `pred_Q`, `pred_VP`
- ▶ Synonyms
 - ▶ `pred_VC`, `pred_V2` and `pred_V` are synonymous for central volume
 - ▶ `pred_VP` and `pred_V3` are synonymous for peripheral volume

\$MAIN [C++]

- ▶ Like \$PK, code that gets called right before solving odes from t_1 to t_2
- ▶ Set initial conditions
- ▶ Derive new variables ... includes covariate modeling
 - ▶ Must type new variables ... usually double or bool
 - ▶ By default, new variables are initialized globally

```
$MAIN  
RESP_0 = KIN/KOUT;  
  
double CLi = exp(log(TVCL) + ETA(1));  
  
double base = 0;  
  
if(NEWIND <= 1) base = 2;
```

ACCESS parameters, compartment amounts, compartment initial conditions all variables and macros

\$ODE [C++]

- ▶ Every compartment **needs** an equation
- ▶ Form: `dxdt_CMT = ratein - rateout;`
- ▶ Use CMT for compartment amounts, parameters, or other variables
- ▶ This block of code gets called repeatedly so be wise
- ▶ May specify new variables here as in \$MAIN; they will be initialized globally

```
$ODE  
dxdt_CENT  = -(CL/VC)*CENT;  
  
double CP = CENT/VC;  
dxdt_RESP = KIN*(1-CP/(IC50+CP)) - KOUT;
```

\$TABLE [C++] and \$CAPTURE [txt]

- ▶ Used to derive variables **after** the solver advances the system
- ▶ Specify additional simulation outputs
 - ▶ Derived variables
 - ▶ Internal variables (like ETA(n) or EPS(n))
 - ▶ Parameters
- ▶ Do not table compartments or ID or TIME
- ▶ \$CAPTURE is convenience wrapper for table statement

```
$TABLE  
table(logRESP) = log(RESP);
```

\$CAPTURE

- Items in \$CAPTURE get translated to `table(name) = name;` in \$TABLE

```
$PARAM FLAG=2, TVCL=1, VC=12
$CMT CENT
$MAIN double CLi = TVCL*exp(ETA(1));
$ODE
double rateout = (CLi/VC)*CENT;
dxdt_CENT = -rateout;
$CAPTURE rateout CLi FLAG
```

- These are equivalent

```
$TABLE table(CP) = CP;
```

```
$CAPTURE CP
```

Example: Simple PK

```
$PROB
PK base model development started 3/12/2016
Kyle Baron

$PARAM CL=1, VC=30

$CMT CENT

$ODE
dxdt_CENT = -(CL/VC)*CENT;

$TABLE table(CP) = CENT/VC;
```

Example: IRM with reduced bioavailability

```
$PARAM CL=1, VC=30, KA=1.1  
KIN=100, KOUT=2, IC50 = 2, FBIO=0.6  
  
$CMT CENT RESP  
  
$MAIN  
RESP_0 = KIN/KOUT;  
_F(1) = FBIO;  
  
$ODE  
double CP = CENT/VC;  
dxdt_CENT = -(CL/VC)*CENT;  
dxdt_RESP = KIN*(1-CP/(IC50+CP)) - KOUT*RESP;  
  
$CAPTURE FBIO
```

\$OMEGA and \$SIGMA [txt]

- ▶ Specify random effect variance/covariance matrices
 - ▶ $\eta_i \sim N(0, \Omega)$
 - ▶ $\varepsilon_j \sim N(0, \Sigma)$
- ▶ NONMEM-style input - lower triangle
- ▶ Multiple \$OMEGA and \$SIGMA are allowed and each may be named
 - ▶ How you enter these blocks defines a “signature” that mrgsolve will enforce later on
- ▶ Diagonal matrix (3x3 in this example)

```
$OMEGA 0.1 0.2 0.3
```


More \$OMEGA and \$SIGMA

- ▶ Options are name=value and must be on different line from the matrix data
- ▶ Block matrix (2x2 in this example)

```
$OMEGA block=TRUE  
0.1 0.06 0.2
```

- ▶ Block matrix with correlations (assumes block=TRUE)

```
$OMEGA cor=TRUE  
0.1 0.67 0.2
```

More \$OMEGA and \$SIGMA

- ▶ Multiple \$OMEGA and \$SIGMA are allowed; each may be named

```
$OMEGA name='PK'
```

```
0 0 0
```

```
$OMEGA name='PD'
```

```
0 0
```

Example: Population PK

```
$PARAM LTVCL=log(1), LTVVC=log(30), WT = 70, PROT=1
```

```
$CMT CENT
```

```
$MAIN
```

```
double CL = exp(LTVCL + 0.75*log(WT/70.0) + ETA(1));
```

```
double VC = exp(LTVVC + *log(WT/70.0) + ETA(2));
```

```
$OMEGA cor=TRUE
```

```
0.2 0.67 0.2
```

```
$SIGMA 0.025 0.1
```

```
$ODE dxdt_CENT = -(CL/VC)*CENT;
```

```
$TABLE
```

```
double eps = PROT==1 ? EPS(1) : EPS(2);
```

```
table(DV) = (CENT/VC)*exp(eps);
```

```
$CAPTURE CL
```

C++ expressions and functions

```
if(a == 2) b = 2;
```

```
if(b <= 2) {
```

```
    c=3;
```

```
} else {
```

```
    c=4;
```

```
}
```

```
double d = pow(base,exponent);
```

```
double e = exp(3);
```

```
double f = fabs(-4);
```

```
double g = sqrt(5);
```

```
double h = log(6);
```

```
double i = log10(7);
```

```
double j = floor(4.2);
```

```
double k = ceil(4.2);
```

\$GLOBAL [C++]

- ▶ Used to define C++ variables or data structures “globally”
 - ▶ Here, “globally” refers to the model shared object
 - ▶ Variables won’t clash with R or mrgsolve internals
- ▶ Also define C++ preprocessor directives here

```
$GLOBAL
```

```
bool cured = false;  
double mult = 0;
```

```
#define g 9.8  
#define plus(a,b) (a+b)
```

```
$GLOBAL
```

```
std::vector<double> a(5);  
double mult(double a, double b) {return a*b;}
```

\$SET [R]

- ▶ Define some settings
 - ▶ Simulation times: start, end, delta, add
 - ▶ Solver settings: hmax, atol, rtol, maxsteps
 - ▶ Compartment requests: request
 - ▶ Time re-scaling: tscale
- ▶ All of these settings can get changed later, but setting them here allows you bind some default settings to the specification file For example:

```
$SET  
end=24*7, delta=1  
hmax=0.1  
tscale = 1/24
```

\$THETA [txt]

```
$THETA
```

```
0.1 0.2 0.3
```

- ▶ Create items in the parameter list with name THETAn
- ▶ A quick way to get THETA-related parameters in to the list without a lot of typing
- ▶ Add a name argument to change the prefix

```
$THETA name="theta"
```

```
1 2 3
```

- ▶ Will insert theta1=1, theta2=2 and theta3=3 into parameter list

\$THETA example

Data items in \$THETA get inserted into \$PARAM, but with an assumed name

```
code <- '  
$THETA 8 9 10  
$PARAM CL=4, VC=5  
'  
mread(code=code, "modspec", proj) %>% param
```

```
.  
.  Model parameters (N=5):  
.  name    value . name    value  
.  CL      4     | THETA3  10  
.  THETA1  8     | VC      5  
.  THETA2  9     | .      .
```


\$FIXED [R]

```
$FIXED
```

```
g = 9.8
```

```
$PARAM
```

```
TVCL = 5
```

- ▶ Functions just like \$PARAM, but values are eternally fixed ... even in R
- ▶ Declared as `const double`
- ▶ Usually only used when there is a large parameter list
 - ▶ Some performance improvement
 - ▶ Makes the parameter list smaller and easier to deal with
- ▶ **ACCESS** R global environment

- ▶ Use \$VCMT to implement an indicator compartment
 - ▶ mrgsolve will write `dxdt_FLAG=0;`
 - ▶ You can dose into or use value in that compartment anywhere in the model
 - ▶ Just for convenience

\$VCMT FLAG

\$ADVAN2 [txt]

- ▶ Including this block tells mrgsolve to implement a one-compartment model with first-order absorption
- ▶ Must include \$CMT with 2 compartments
 - ▶ First name is the dosing compartment and second is the central compartment
 - ▶ But you decide on the names

```
$PARAM CL=1, VC=20, KA=1.1
```

```
$CMT DEPOT CENT
```

```
$MAIN
```

```
double CLi = exp(log(CL) + ETA(1));
```

```
pred_CL = CLi;
```

```
pred_VC = VC;
```

```
pred_KA = KA;
```

\$ADVAN4 [txt]

- ▶ Including this block tells mrgsolve to implement a two-compartment model with first-order absorption
- ▶ Must include CMT with 3 compartments
 - ▶ First name is dosing, second is central, third is peripheral
 - ▶ But you decide on the names

```
$PARAM CL=1, VC=20, KA=1.1
```

```
VP = 120 Q=5
```

```
$CMT DEPOT CENT PERIPH
```

```
$MAIN
```

```
double CLi = exp(log(CL) + ETA(1));
```

```
pred_CL = CLi;
```

```
pred_VC = VC;
```

```
pred_KA = KA;
```