mrgsolve: Simulate from ODE-Based Models

Get Started Package Vignette

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mrgsolve is an R package maintained under the auspices of Metrum Research Group that facilitates simulation from models based on systems of ordinary differential equations (ODE) that are typically employed for understanding pharmacokinetics, pharmacodynamics, and systems biology and pharmacology. mrgsolve consists of computer code written in the R and C++ languages, providing an interface to a C++ translation of the lsoda differential equation solver.

This vignette will show you how to get started using mrgsolve.

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This vignette introduces the mrgsolve workflow. First, load the package along with any other helper packages we need for this vignette.

```
library(mrgsolve)
library(dplyr)
```

1 Get started

To get started with mrgsolve, try using the built in model library like this

mrgsolve 1 GET STARTED

```
mod <- modlib("pk1", delta = 0.1)

out <- mrgsim(mod, events = ev(amt = 100))

out</pre>
```

```
Model:
        pk1
Dim:
        242 \times 5
Time:
        0 to 24
ID:
        1
    ID time
                 ΕV
                                CP
                      CENT
     1
        0.0
              0.00
                    0.000 0.0000
1:
                     0.000 0.0000
2:
        0.0 100.00
        0.1
             90.48
                    9.492 0.4746
4:
        0.2
             81.87 18.034 0.9017
5:
        0.3
             74.08 25.715 1.2858
             67.03 32.619 1.6309
6:
        0.4
7:
        0.5
             60.65 38.819 1.9409
       0.6 54.88 44.383 2.2191
```

```
plot(out, "CP")
```

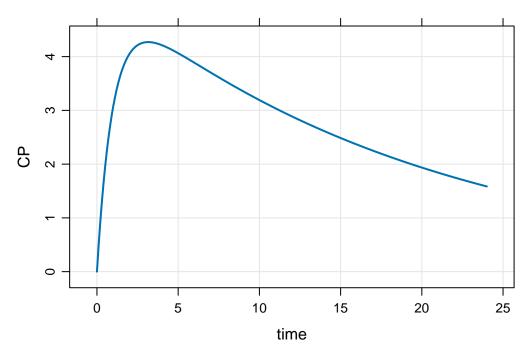


Figure 1: Simple simulation of a single dose

That was a really simple simulation where we used an event object to initiate a dose into a one-compartment model. See how the plot() method allows us to quickly visualize what happend in the simulation.

2 Event objects

Event objects help you implement dosing events with a lightweight, easy to compose syntax. You construct them with the ev() function. So I can make an object for a single 100 mg bolus dose

```
ev(amt = 100)
```

Events:

```
time amt cmt evid
1 0 100 1 1
```

or we can code a series of intermittent infusions

```
ev(amt = 100, rate = 50, ii = 24, addl = 3)
```

Events:

```
time amt rate ii addl cmt evid 1 \quad 0 \quad 100 \quad 50 \quad 24 \quad 3 \quad 1 \quad 1
```