Conditional Dosing Pharmacometric Example

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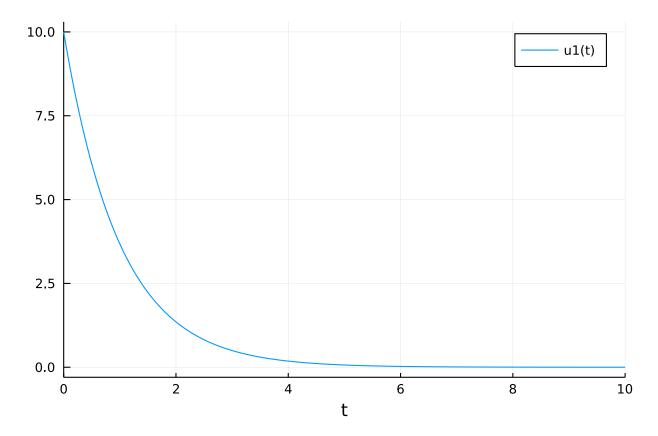
In this example we will show how to model a conditional dosing using the DiscreteCallbacks. The problem is as follows. The patient has a drug A(t) in their system. The concentration of the drug is given as C(t)=A(t)/V for some volume constant V. At t=4, the patient goes to the clinic and is checked. If the concentration of the drug in their body is below 4, then they will receive a new dose.

For our model, we will use the simple decay equation. We will write this in the in-place form to make it easy to extend to more complicated examples:

```
using DifferentialEquations
function f(du,u,p,t)
    du[1] = -u[1]
end
u0 = [10.0]
const V = 1
prob = ODEProblem(f,u0,(0.0,10.0))

ODEProblem with uType Vector{Float64} and tType Float64. In-place: true timespan: (0.0, 10.0)
u0: 1-element Vector{Float64}:
10.0

Let's see what the solution looks like without any events.
sol = solve(prob,Tsit5())
using Plots; gr()
plot(sol)
```



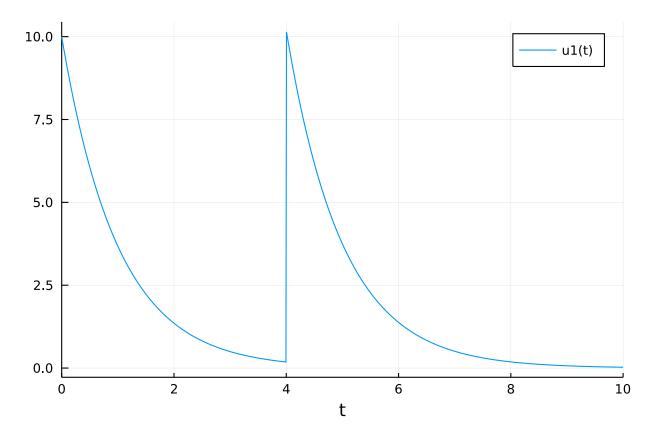
We see that at time t=4, the patient should receive a dose. Let's code up that event. We need to check at t=4 if the concentration u[1]/4 is <4, and if so, add 10 to u[1]. We do this with the following:

```
condition(u,t,integrator) = t==4 && u[1]/V<4
affect!(integrator) = integrator.u[1] += 10
cb = DiscreteCallback(condition,affect!)</pre>
```

DiffEqBase.DiscreteCallback{typeof(Main.##WeaveSandBox#5436.condition), typeof(Main.##WeaveSandBox#5436.affect!), typeof(DiffEqBase.INITIALIZE_DEFAULT), typeof(DiffEqBase.FINALIZE_DEFAULT)}(Main.##WeaveSandBox#5436.condition, Main.##WeaveSandBox#5436.affect!, DiffEqBase.INITIALIZE_DEFAULT, DiffEqBase.FINALIZE_DEFAULT, Bool[1, 1])

Now we will give this callback to the solver, and tell it to stop at t=4 so that way the condition can be checked:

```
sol = solve(prob,Tsit5(),tstops=[4.0],callback=cb)
using Plots; gr()
plot(sol)
```



Let's show that it actually added 10 instead of setting the value to 10. We could have set the value using affect!(integrator) = integrator.u[1] = 10

```
println(sol(4.00000))
println(sol(4.000000000001))

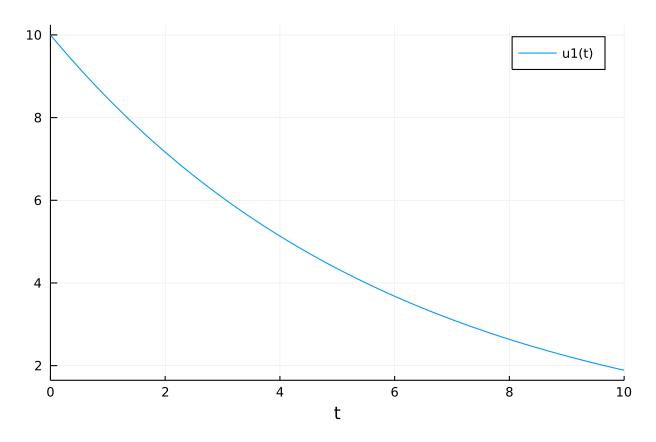
[0.18316389222022983]
[10.183163892210045]
```

Now let's model a patient whose decay rate for the drug is lower:

```
function f(du,u,p,t)
    du[1] = -u[1]/6
end
u0 = [10.0]
const V = 1
prob = ODEProblem(f,u0,(0.0,10.0))

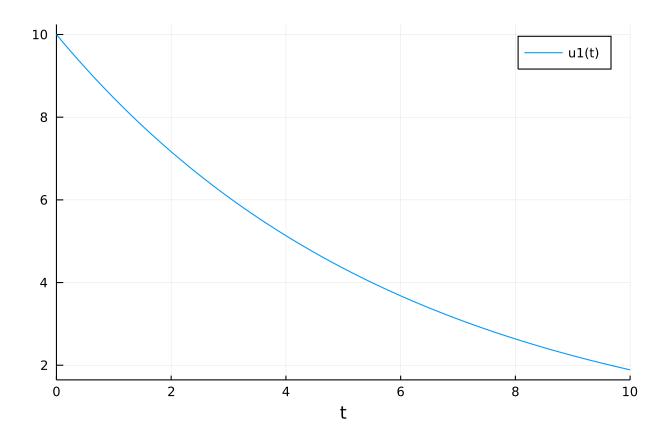
ODEProblem with uType Vector{Float64} and tType Float64. In-place: true timespan: (0.0, 10.0)
u0: 1-element Vector{Float64}:
10.0

sol = solve(prob,Tsit5())
using Plots; gr()
plot(sol)
```



Under the same criteria, with the same event, this patient will not receive a second dose:

```
sol = solve(prob,Tsit5(),tstops=[4.0],callback=cb)
using Plots; gr()
plot(sol)
```



0.1 Appendix

using SciMLTutorials

These tutorials are a part of the SciMLTutorials.jl repository, found at: https://github.com/SciML/SciMLFor more information on high-performance scientific machine learning, check out the SciML Open Source Software Organization https://sciml.ai.

To locally run this tutorial, do the following commands:

```
SciMLTutorials.weave_file("tutorials/models","02-conditional_dosing.jmd")

Computer Information:

Julia Version 1.6.2

Commit 1b93d53fc4 (2021-07-14 15:36 UTC)

Platform Info:
    OS: Linux (x86_64-pc-linux-gnu)
    CPU: AMD EPYC 7502 32-Core Processor
    WORD_SIZE: 64
    LIBM: libopenlibm
    LLVM: libLLVM-11.0.1 (ORCJIT, znver2)

Environment:
    JULIA_DEPOT_PATH = /root/.cache/julia-buildkite-plugin/depots/a6029d3a-f78b-41ea-bc9
    JULIA_NUM_THREADS = 16
```

Package Information:

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[f3b72e0c] DiffEqDevTools v2.27.2
[055956cb] DiffEqPhysics v3.9.0
\hbox{\tt [Oc46a032] Differential Equations v6.17.1}
[31c24e10] Distributions v0.24.18
[587475ba] Flux v0.12.1
[f6369f11] ForwardDiff v0.10.18
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```

And the full manifest:

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