

Simplest COVID model

```
# Initial conditions of the state variables (D=deceased)
yini <- c(S = 17.5e6, I = 1e3, R = 0, D = 0) # number of people

# Model parameters
pars <- c(
  b = 0.00000002, # [/ind/d] infection rate constant
  g = 0.07,       # [/d] recovery rate constant
  m = 0.007       # [/d] mortality rate constant
)

# Model function: calculates time-derivatives and other output
SIRmodel <- function(t, state, pars) {
  # t: time, state: state variables, pars: model parameters
  with (as.list(c(state, pars)), {

    # rate expressions [ind/d]
    Infection <- b * I * S # infection rate
    Recovery <- g * I      # recovery rate
    Mortality <- m * I     # mortality rate

    # Time-derivatives: dC/dt = production - consumption [ind/d]
    dSdt <- -Infection
    dIdt <- Infection - Recovery - Mortality
    dRdt <- Recovery
    dDdt <- Mortality

    # return time-derivatives and ordinary variables as a list
    list(c(dSdt, dIdt, dRdt, dDdt), # vector with derivatives
         # (the same order as state variables!)
         Infection = Infection, # other output
         Mortality = Mortality)
  })
}
```

```
require(deSolve) # package with integration methods
# vector of output times
outtimes <- seq(from = 1, to = 100, length.out = 100)
# ode integrates the model
out <- ode(y=yini, parms=pars, func=SIRmodel, times=outtimes)
# plot the model output
plot(out)
```

Enhanced COVID model

(a) →

```
# define forcing function for the social interactions parameter
# sint = 1: no restrictions, sint = 0.1: restricted to 10% of normal
sIntDATA <- data.frame(time = c(0, 29, 30, 69, 70, 100),
                        sInt = c(1, 1, 0.1, 0.1, 1, 1))
sIntFUN <- approxfun(x=sIntDATA)

# Model function: calculates time-derivatives and other output
SIRmodel2 <- function(t, state, pars, sIntDyn) { # (b)
  # t: time, state: state variables, pars: model parameters
  with (as.list(c(state, pars)), {

    # social interactions parameter sint is determined externally
    sInt <- sIntDyn(t)

    # rate expressions [ind/d]
    Infection <- sInt * b * I * S # infection rate
    Recovery <- g * I              # recovery rate
    Mortality <- m * I             # mortality rate

    # Time-derivatives: dC/dt = production - consumption [ind/d]
    dSdt <- -Infection
    dIdt <- Infection - Recovery - Mortality
    dRdt <- Recovery
    dDdt <- Mortality

    # return time-derivatives and ordinary variables as a list
    list(c(dSdt, dIdt, dRdt, dDdt), # vector with derivatives
         # (the same order as state variables!)
         Infection = Infection,      # other output
         Mortality = Mortality)
  })
}
```

(c) →

```
require(deSolve)
# ode integrates the model
out2 <- ode(y=yini, parms=pars, func=SIRmodel2, times=outtimes,
            sIntDyn=sIntFUN) # forcing function included
# plot the model output
plot(out, out2)
```